

1343

1344

NT2RP4001725	18.61	25.02	18.94	1	1
NT2RP4001726	66.49	89.94	73.84	1.35	1.11
NT2RP4001730	70.16	63.61	47.29	0.91	0.67
NT2RP4001739	42.06	51.3	49.42	1.22	1.17
NT2RP4001741	180.84	250.66	135.6	1.39	0.75
NT2RP4001753	72.78	104.66	52.65	1.44	0.72
NT2RP4001760	51	52.44	44.59	1.03	0.87
NT2RP4001787	4461.9	5125.17	3639.25	1.15	0.82
NT2RP4001790	46.27	54.59	40.79	1.18	0.88
NT2RP4001795	154.14	189.71	153.64	1.23	1
NT2RP4001803	50.74	51.85	46.05	1.02	0.91
NT2RP4001805	32.65	36.8	29.6	1	1
NT2RP4001809	191.23	213.41	137.81	1.12	0.72
NT2RP4001817	112.92	162.95	120.23	1.44	1.06
NT2RP4001822	65.32	70.78	65.97	1.08	1.01
NT2RP4001823	21.06	22.44	21.32	1	1
NT2RP4001827	102.8	112	95.96	1.09	0.93
NT2RP4001828	431.57	583.31	473.45	1.35	1.1
NT2RP4001836	146.86	202.66	148.56	1.38	1.01
NT2RP4001838	29.91	43.1	37.71	1.08	1
NT2RP4001841	1549.1	1954.07	1139.77	1.26	0.74
NT2RP4001849	33.35	29.78	25.37	1	1
NT2RP4001861	736.02	720.46	638.48	0.98	0.87
NT2RP4001877	93.22	103.29	93.49	1.11	1
NT2RP4001879	49.34	70.19	54.7	1.42	1.11
NT2RP4001889	103.01	187.69	165.9	1.82	1.61
NT2RP4001893	27.15	40.29	34.35	1.01	1
NT2RP4001896	37.41	97.88	61.56	2.45	1.54
NT2RP4001898	183.3	215.58	185.12	1.18	1.01
NT2RP4001901	60.98	80.96	58.59	1.33	0.96
NT2RP4001910	408.91	559.44	327.12	1.37	0.8
NT2RP4001925	59.81	72.23	51.93	1.21	0.87
NT2RP4001926	46.24	66.65	60.62	1.44	1.31
NT2RP4001927	42.17	51.54	45.85	1.22	1.09
NT2RP4001931	54.82	81.7	62.6	1.49	1.14
NT2RP4001933	181.43	229.76	269.85	1.27	1.49
NT2RP4001938	89.48	111.84	97.72	1.25	1.09
NT2RP4001942	157.21	178.4	186.22	1.13	1.18
NT2RP4001945	23.76	21.74	26.27	1	1
NT2RP4001946	34.33	46.5	40.43	1.16	1.01
NT2RP4001947	18.29	28.67	26.12	1	1
NT2RP4001950	23.81	28.29	30.45	1	1
NT2RP4001953	76.32	134.32	95.15	1.76	1.25
NT2RP4001966	14.53	24.06	18.92	1	1
NT2RP4001970	32.46	66.99	45.62	1.67	1.14
NT2RP4001975	93.45	117.02	108.65	1.25	1.16
NT2RP4001988	103.89	188.4	157.9	1.81	1.52
NT2RP4001996	47.79	87.05	68.23	1.82	1.43
NT2RP4002014	48.78	113.53	89.64	2.33	1.84
NT2RP4002018	50.95	67.62	72.16	1.33	1.42

(674)

特開 2002-191363

1345

1346

NT2RP4002035	38.64	83.62	57.12	2.09	1.43
NT2RP4002043	50.55	74.39	61.28	1.47	1.21
NT2RP4002046	113.68	146.69	103.17	1.29	0.91
NT2RP4002047	96.67	132.21	84.39	1.37	0.87
NT2RP4002052	47.17	63.23	54.34	1.34	1.15
NT2RP4002056	350.57	455.24	380.17	1.3	1.08
NT2RP4002057	61.49	117.62	116.72	1.91	1.9
NT2RP4002058	25.72	47.72	52.24	1.19	1.31
NT2RP4002064	19.42	33.93	26.12	1	1
NT2RP4002071	44.49	82.96	52.47	1.86	1.18
NT2RP4002075	28.6	36.73	33.16	1	1
NT2RP4002078	92.71	143.87	108.63	1.55	1.17
NT2RP4002081	236.65	249.16	249.24	1.05	1.05
NT2RP4002083	17.72	23.02	27.12	1	1
NT2RP4002099	18.6	29.18	27.89	1	1
NT2RP4002106	54.11	107.08	119	1.98	2.2
NT2RP4002111	101.96	200.72	148.95	1.97	1.46
NT2RP4002112	74.68	68.41	53.51	0.92	0.72
NT2RP4002116	379.37	384.55	382.82	1.01	1.01
NT2RP4002122	61.29	75.89	55.81	1.24	0.91
NT2RP4002126	66.06	115.78	77.24	1.75	1.17
NT2RP4002133	124.26	188.7	156.52	1.52	1.26
NT2RP4002136	59.72	83.01	65.67	1.39	1.1
NT2RP4002139	196.25	298.89	317.57	1.52	1.62
NT2RP4002174	47.12	56.4	49.37	1.2	1.05
NT2RP4002185	60.33	83.37	80.31	1.38	1.33
NT2RP4002186	122.78	157.98	101.33	1.29	0.83
NT2RP4002187	105.68	137.58	144.38	1.3	1.37
NT2RP4002188	70.98	106.05	83.05	1.49	1.17
NT2RP4002199	31.14	47.63	46.66	1.19	1.17
NT2RP4002206	39.97	38.91	31.55	1	1
NT2RP4002210	26.97	23.02	24.96	1	1
NT2RP4002222	38.72	58.99	42.3	1.47	1.06
NT2RP4002241	81.81	88.06	74.33	1.08	0.91
NT2RP4002248	85.52	108.46	83.74	1.27	0.98
NT2RP4002250	6.36	20.54	11.35	1	1
NT2RP4002259	29.41	42.66	31.34	1.07	1
NT2RP4002268	159.71	202.23	161.92	1.27	1.01
NT2RP4002288	181.37	257.54	204.61	1.42	1.13
NT2RP4002290	56.9	76.7	71.74	1.35	1.26
NT2RP4002298	53.47	84.08	46.67	1.57	0.87
NT2RP4002306	53.35	84.82	74.88	1.59	1.4
NT2RP4002308	42.24	41.6	33.19	0.98	0.95
NT2RP4002336	68.71	90.07	67.38	1.31	0.98
NT2RP4002340	19.09	13.39	14.64	1	1
NT2RP4002361	40.9	56.53	37.12	1.38	0.98
NT2RP4002367	29.1	25.75	23.97	1	1
NT2RP4002368	94.98	113.04	103.33	1.19	1.09
NT2RP4002377	289.2	315.73	317.74	1.09	1.1
NT2RP4002408	30.29	33.42	32.09	1	1

(675)

特開 2002-191363

1347

1348

NT2RP4002425	43.47	72.09	45.88	1.66	1.06
NT2RP4002432	1549.3	2074.3	1253.27	1.34	0.81
NT2RP4002447	99.1	100.15	81.33	1.01	0.82
NT2RP4002451	55.07	53.75	50.58	0.98	0.92
NT2RP4002461	283.54	296.78	229.24	1.05	0.81
NT2RP4002486	52.21	62.8	51.82	1.2	0.99
NT2RP4002517	86.77	89.86	85.96	1.04	0.99
NT2RP4002556	56.11	73.77	77.16	1.31	1.38
NT2RP4002569	96.58	96.6	81.78	1	0.85
NT2RP4002587	59.69	46.93	47.79	0.79	0.8
NT2RP4002591	59.54	66.69	61.58	1.12	1.03
NT2RP4002607	65.45	67.05	57.12	1.02	0.87
NT2RP4002627	1154.4	1435.6	1039.2	1.24	0.9
NT2RP4002628	89.25	115.91	86.23	1.3	0.97
NT2RP4002630	139.5	192.58	143.27	1.38	1.03
NT2RP4002639	949.82	913.58	777.44	0.96	0.82
NT2RP4002641	57.68	78.81	58.54	1.37	1.01
NT2RP4002658	690.44	903.21	768.32	1.31	1.11
NT2RP4002669	55.37	58.95	55.54	1.06	1
NT2RP4002677	65.98	122.24	115.69	1.85	1.75
NT2RP4002715	160.17	197.94	157.66	1.24	0.98
NT2RP4002750	25.5	31.12	32	1	1
NT2RP4002784	79.32	110.67	79.11	1.4	1
NT2RP4002791	31.48	81.22	60.98	2.03	1.52
NT2RP4002811	30.15	37.16	31.27	1	1
NT2RP4002830	78.6	93.65	83.44	1.19	1.06
NT2RP4002832	26.95	34.45	33.04	1	1
NT2RP4002850	105.03	134.98	121.72	1.29	1.16
NT2RP4002874	14.39	16.4	17.69	1	1
NT2RP4002884	248.98	397.43	368.37	1.6	1.48
NT2RP4002888	22.67	39.71	35.05	1	1
NT2RP4002891	78.58	105.26	101.71	1.34	1.29
NT2RP4002894	157.28	186.39	182.82	1.19	1.16
NT2RP4002896	62.43	72.07	61.47	1.15	0.98
NT2RP4002905	35.39	37.85	36.81	1	1
NT2RP4002907	60.03	78.35	74.11	1.31	1.23
NT2RP5003459	2814.9	3301.39	3101.17	1.17	1.1
NT2RP5003461	46.24	61.77	44.54	1.34	0.96
NT2RP5003471	2383.9	4003.5	2319.1	1.68	0.97
NT2RP5003477	14.46	21.09	22.54	1	1
NT2RP5003487	4297.0	5054.14	3381.8	1.18	0.79
NT2RP5003492	31.05	42.59	32.16	1.06	1
NT2RP5003500	28.6	32.69	26.16	1	1
NT2RP5003506	73.76	122.61	102.08	1.66	1.38
NT2RP5003512	15.54	30.94	28.8	1	1
NT2RP5003522	18.03	33.33	26.71	1	1
NT2RP5003524	17.34	31.72	29.03	1	1
NT2RP5003527	349.06	483.98	393.69	1.39	1.13
NT2RP5003531	75.41	76.61	66.44	1.02	0.88
NT2RP5003534	52.83	100.86	54.98	1.91	1.04

(676)

特開 2002-191363

1349

1350

NT2RP6000020	254.97	316.92	257.92	1.24	1.01
NT2RP6000022	28.24	42.02	41.88	1.05	1.05
NT2RP6000050	21.19	34.01	39.08	1	1
NT2RP6000063	24.89	31.66	29.69	1	1
NT2RP6000074	27.19	40.18	36.98	1	1
NT2RP6000083	68.33	91.18	64.08	1.33	0.94
NT2RP6000100	52.92	50.18	45.21	0.95	0.85
NT2RP6000123	34.42	63.3	47.29	1.58	1.18
NT2RP6000129	40.69	50.73	39.14	1.25	0.98
NT2RP6000147	87.35	87.01	84.55	1	0.97
NT2RP6000163	28.6	34.59	32.31	1	1
NT2RP6000181	69.12	113.55	90.59	1.64	1.31
NT2RP6000182	42.14	68.54	51.62	1.63	1.22
OVARC1000001	49.41	91.19	69.76	1.85	1.41
OVARC1000003	102.25	87.38	70.78	0.85	0.69
OVARC1000004	2892.4	2673.1	1762.66	0.92	0.61
OVARC1000006	42.54	69.38	47.66	1.63	1.12
OVARC1000013	63.38	78.08	67.4	1.23	1.06
OVARC1000014	42.8	52.28	43.62	1.22	1.02
OVARC1000017	36.69	47.86	48.03	1.2	1.2
OVARC1000026	320.47	405.43	316.62	1.27	0.99
OVARC1000035	113.44	134.98	108.81	1.19	0.96
OVARC1000037	134.37	144.88	164.84	1.08	1.23
OVARC1000058	73.52	87.2	80.14	1.19	1.09
OVARC1000060	64.66	84.81	57.28	1.31	0.89
OVARC1000068	38.02	38.3	43.24	1	1.08
OVARC1000069	40.42	57.23	49.59	1.42	1.23
OVARC1000071	32.24	33.2	29.25	1	1
OVARC1000075	2412.8	3591.21	2207.17	1.49	0.91
OVARC1000083	117.97	119.13	69.75	1.01	0.59
OVARC1000085	792.51	498.22	687.24	0.63	0.87
OVARC1000086	69.36	75.22	72.13	1.08	1.04
OVARC1000087	24.26	30.25	33.69	1	1
OVARC1000090	123.62	113.17	136.64	0.92	1.11
OVARC1000091	41.32	40.24	43.17	0.97	1.04
OVARC1000092	59.92	45.23	66.35	0.75	1.11
OVARC1000105	193.62	211.28	213.54	1.09	1.1
OVARC1000106	207.32	203.69	194.81	0.98	0.94
OVARC1000109	95.88	83.7	80.21	0.87	0.84
OVARC1000113	119.41	112.39	127.87	0.94	1.07
OVARC1000114	92.45	81.84	102.21	0.89	1.11
OVARC1000133	40.28	48	69.51	1.19	1.73
OVARC1000137	68.99	58.7	79.44	0.85	1.15
OVARC1000139	1597.0	1399.58	1239.81	0.88	0.78
OVARC1000145	18.41	25.65	25.84	1	1
OVARC1000148	80.77	84.79	78.69	1.05	0.97
OVARC1000151	38.7	34.81	30.58	1	1
OVARC1000157	126.85	138.69	121.57	1.09	0.96
OVARC1000162	31.19	30.47	41.58	1	1.04
OVARC1000168	60.64	84.53	66.09	1.39	1.09

(677)

特開 2002-191363

1351

1352

OVARC1000169	593.72	714.83	894.21	1.2	1.51
OVARC1000178	39.59	38.33	28.67	1	1
OVARC1000182	20.83	21.3	22.61	1	1
OVARC1000186	49.43	45.22	42.58	0.91	0.86
OVARC1000188	29.19	26.83	21.31	1	1
OVARC1000191	26.54	29.18	21.28	1	1
OVARC1000198	62.71	85.36	58.69	1.36	0.94
OVARC1000208	272.18	324.49	263.93	1.19	0.97
OVARC1000209	141.71	150.62	119.37	1.06	0.84
OVARC1000212	47.05	49.04	39.68	1.04	0.85
OVARC1000216	17.93	18.99	19.61	1	1
OVARC1000240	103.46	101.08	71.71	0.98	0.69
OVARC1000241	47.76	37.11	42.61	0.84	0.89
OVARC1000249	34.16	38.99	36.88	1	1
OVARC1000254	1580.5	2172.35	1417.43	1.37	0.9
OVARC1000255	26.25	24.08	22.57	1	1
OVARC1000267	158.61	196.13	154.6	1.24	0.97
OVARC1000275	27.34	25.99	28.89	1	1
OVARC1000287	62.95	64.26	84.75	1.02	1.35
OVARC1000288	64.73	68.75	45.65	1.06	0.71
OVARC1000298	79.52	94.53	100.14	1.19	1.26
OVARC1000302	30.55	38.34	29.12	1	1
OVARC1000304	37.61	40.63	42.47	1.02	1.06
OVARC1000307	36.06	33.36	41.11	1	1.03
OVARC1000309	29.18	27.03	22.21	1	1
OVARC1000312	31.23	38.59	29.77	1	1
OVARC1000313	102.54	157.6	184.77	1.54	1.8
OVARC1000321	152.01	148.5	138.68	0.98	0.91
OVARC1000326	27.03	34.02	23.17	1	1
OVARC1000327	29.37	30.83	26.7	1	1
OVARC1000331	47.64	62.99	51.7	1.32	1.09
OVARC1000335	27.77	28.39	32.55	1	1
OVARC1000347	48.79	55.07	37.14	1.13	0.82
OVARC1000348	17.11	23.34	19.96	1	1
OVARC1000363	42.47	45.27	38.6	1.07	0.94
OVARC1000377	29.55	27.57	31.2	1	1
OVARC1000382	31.12	37.4	34.78	1	1
OVARC1000384	37.91	42.89	31.82	1.07	1
OVARC1000401	27.05	26.01	18.81	1	1
OVARC1000406	1432.1	1992.66	2667.11	1.39	1.86
OVARC1000407	55.42	62.17	49.83	1.12	0.9
OVARC1000408	521.33	474.3	454.23	0.91	0.87
OVARC1000410	57.23	48.05	63.58	0.84	1.11
OVARC1000411	36.21	41.64	34.64	1.04	1
OVARC1000414	102.21	105.16	89.37	1.03	0.87
OVARC1000420	49.94	53.08	44.19	1.06	0.88
OVARC1000421	47.02	45.9	44.27	0.98	0.94
OVARC1000427	3330.4	3323.93	2479.35	1	0.74
OVARC1000431	163.82	212.48	204.37	1.3	1.25
OVARC1000437	52.04	48.36	47.4	0.93	0.91

(678)

特開 2002-191363

1353

1354

OVARC1000439	113.64	76.62	106.88	0.67	0.94
OVARC1000440	71.11	74.54	56.39	1.05	0.79
OVARC1000442	119.54	121.73	109.35	1.02	0.91
OVARC1000443	65.66	52.35	38.24	0.8	0.61
OVARC1000461	29.17	37.12	25	1	1
OVARC1000465	34.43	34.38	43.55	1	1.09
OVARC1000466	46.92	37.83	43.09	0.85	0.92
OVARC1000467	38.38	36.82	38.22	1	1
OVARC1000470	62.44	77.76	68.31	1.25	1.09
OVARC1000473	47.79	41.99	31.39	0.88	0.84
OVARC1000479	83.4	72.71	77.11	0.87	0.92
OVARC1000484	116.94	161.5	98.17	1.38	0.84
OVARC1000486	54.85	62.16	47.76	1.13	0.87
OVARC1000496	18.29	28.12	16.71	1	1
OVARC1000520	24.36	22.81	23.25	1	1
OVARC1000522	64	68.45	54.56	1.07	0.85
OVARC1000526	102.56	106.43	75.86	1.04	0.74
OVARC1000529	54.69	60.42	41.68	1.1	0.76
OVARC1000533	80.21	81.45	73.46	1.02	0.92
OVARC1000543	36.03	34.69	22.16	1	1
OVARC1000550	33.22	39.9	37.09	1	1
OVARC1000553	78.24	110.76	91.37	1.42	1.17
OVARC1000556	68.77	93.52	73.85	1.36	1.07
OVARC1000557	28.95	42.48	33.83	1.06	1
OVARC1000561	257.61	351.52	287.81	1.36	1.12
OVARC1000564	139.65	110.7	130.87	0.79	0.94
OVARC1000573	68.41	59.91	62.83	0.88	0.92
OVARC1000576	2089.9	1861.86	1692.32	0.89	0.81
OVARC1000578	43.7	34.85	48.46	0.92	1.11
OVARC1000581	20.13	21.57	21.97	1	1
OVARC1000586	360.06	393.98	421.49	1.09	1.17
OVARC1000588	27.3	24.22	27.51	1	1
OVARC1000605	48.33	40.91	46.84	0.85	0.97
OVARC1000622	225.87	248.61	210.75	1.1	0.93
OVARC1000636	31.45	27.78	33.66	1	1
OVARC1000640	51.97	54.12	101.55	1.04	1.95
OVARC1000649	1164.0	1285.41	980.59	1.1	0.84
OVARC1000661	50.99	49.55	47.02	0.97	0.92
OVARC1000677	132.42	141.8	167.49	1.07	1.26
OVARC1000678	54.44	54.52	57.5	1	1.06
OVARC1000679	51.06	54.02	40.27	1.06	0.79
OVARC1000681	38.25	27.56	28.08	1	1
OVARC1000682	341.04	412.78	251.86	1.21	0.74
OVARC1000689	99.5	105.11	89.74	1.06	0.9
OVARC1000700	58.34	75.6	60.57	1.3	1.04
OVARC1000703	73.66	74.87	53.93	1.02	0.73
OVARC1000722	715.27	635.83	473.66	0.89	0.66
OVARC1000726	38.23	28.7	25.99	1	1
OVARC1000727	48.09	50.07	40.26	1.04	0.84
OVARC1000730	73.44	74.87	80.56	1.02	1.1

(679)

特開 2002-191363

1355

1356

OVARC1000741	73.45	80.84	89.53	1.1	1.22
OVARC1000746	26.64	30	28.58	1	1
OVARC1000764	88.26	97.96	68.38	1.11	0.77
OVARC1000769	101.98	141.42	90.47	1.39	0.89
OVARC1000771	40.54	47.82	40.96	1.18	1.01
OVARC1000773	255.51	361.9	494.36	1.42	1.93
OVARC1000775	115.23	107.57	86.04	0.93	0.75
OVARC1000778	55.18	60.68	47.93	1.1	0.87
OVARC1000779	15.03	16.56	13.23	1	1
OVARC1000781	33.98	45.89	37.86	1.15	1
OVARC1000787	35.99	33.77	27.66	1	1
OVARC1000789	112.59	74.08	135.19	0.66	1.2
OVARC1000800	85.36	118.32	95.97	1.39	1.12
OVARC1000802	39.52	39.36	32.48	1	1
OVARC1000810	119.13	160.74	97.12	1.35	0.82
OVARC1000811	39.94	51.48	34.85	1.29	1
OVARC1000814	174.62	239.67	149.65	1.37	0.86
OVARC1000816	47.23	66.84	64.23	1.42	1.36
OVARC1000817	22.37	20.91	18.09	1	1
OVARC1000834	26.16	25	26.88	1	1
OVARC1000846	73.93	93.47	72.32	1.26	0.98
OVARC1000850	31.18	37.53	39.12	1	1
OVARC1000853	171.25	177.98	145.03	1.04	0.85
OVARC1000862	37.1	42.09	43.93	1.05	1.1
OVARC1000873	60.05	84.54	94.75	1.41	1.58
OVARC1000875	36.96	37.79	30.93	1	1
OVARC1000876	32.05	23.99	36.5	1	1
OVARC1000883	49.62	61.62	63.13	1.24	1.27
OVARC1000885	34.95	40.51	53.3	1.01	1.33
OVARC1000886	44.21	43.73	64.9	0.99	1.47
OVARC1000890	1660.8	1784.57	1371.74	1.07	0.83
OVARC1000891	40.25	41.98	35.62	1.04	0.99
OVARC1000897	25.06	19.74	16.49	1	1
OVARC1000912	59.56	69.38	47.1	1.16	0.79
OVARC1000914	40.41	41.76	36	1.03	0.99
OVARC1000915	49.67	62.61	41.44	1.26	0.83
OVARC1000916	47.9	43.94	45.57	0.92	0.95
OVARC1000924	29.28	34.41	30.96	1	1
OVARC1000928	35.01	25.29	28.53	1	1
OVARC1000936	34.94	26.94	36.31	1	1
OVARC1000937	52.14	62.32	40.83	1.2	0.78
OVARC1000945	27.17	35.2	26.08	1	1
OVARC1000948	19.22	16.97	17.07	1	1
OVARC1000956	65.96	70.99	47.84	1.08	0.73
OVARC1000959	52.49	59.37	43.22	1.13	0.82
OVARC1000960	280.28	287.82	206.56	1.03	0.74
OVARC1000964	2736	2613.45	1992.33	0.96	0.73
OVARC1000971	38.12	24.33	25.94	1	1
OVARC1000975	690.45	651.91	484.63	0.94	0.7
OVARC1000976	29.45	24.47	22.17	1	1

(680)

特開 2002-191363

1357

1358

OVARC1000981	83.75	91.8	88.53	1.1	1.06
OVARC1000982	37.13	52.8	53.82	1.32	1.35
OVARC1000984	30.78	45.59	31.16	1.14	1
OVARC1000995	92.52	90.12	82.09	0.97	0.89
OVARC1000996	43.55	41.13	31.11	0.94	0.92
OVARC1000999	190.43	223.74	180.73	1.17	0.95
OVARC1001000	124.54	163.65	103.45	1.31	0.83
OVARC1001004	20.37	19.73	11.87	1	1
OVARC1001010	28.12	28.13	20.21	1	1
OVARC1001011	53.64	75.49	55.27	1.41	1.03
OVARC1001030	785.13	981.64	1153.75	1.25	1.47
OVARC1001032	39.89	55.5	33.14	1.39	1
OVARC1001034	32.15	37.76	26.08	1	1
OVARC1001038	57.97	61.91	47.15	1.07	0.81
OVARC1001040	161.02	206.83	134.77	1.28	0.84
OVARC1001041	89.09	128.8	96.87	1.45	1.09
OVARC1001044	26.96	24.74	22.65	1	1
OVARC1001049	102.73	174.76	131.25	1.7	1.28
OVARC1001051	824.36	715.74	791.3	0.87	0.96
OVARC1001054	22.98	27.27	27.9	1	1
OVARC1001055	53.59	60.1	38.64	1.12	0.75
OVARC1001062	59.15	51.97	63.58	0.88	1.07
OVARC1001065	428.98	431.9	407.39	1.01	0.95
OVARC1001068	34.9	33.41	45.49	1	1.14
OVARC1001072	26.92	24.25	32.9	1	1
OVARC1001073	36.02	34.51	45.07	1	1.13
OVARC1001074	25.32	30.81	29.07	1	1
OVARC1001078	39.03	51.19	54.45	1.28	1.36
OVARC1001085	47.8	50.43	44.75	1.06	0.94
OVARC1001086	45.34	42.19	44.01	0.93	0.97
OVARC1001091	2172.5	1902.95	1634.39	0.88	0.75
OVARC1001092	41.59	40.51	44.8	0.97	1.08
OVARC1001104	43.01	47.57	49.82	1.11	1.16
OVARC1001107	313.26	304.53	294.7	0.97	0.94
OVARC1001113	24.12	21.01	23.72	1	1
OVARC1001117	73.35	65.99	75.97	0.9	1.04
OVARC1001118	88.32	80.99	78.51	0.92	0.89
OVARC1001125	66.79	70.91	67.69	1.06	1.01
OVARC1001129	39.94	40.36	41.27	1.01	1.03
OVARC1001132	50.72	51.63	68.85	1.02	1.36
OVARC1001138	229.26	273.54	263.88	1.19	1.15
OVARC1001141	50.72	57.25	57.84	1.13	1.14
OVARC1001154	776.93	774.37	672.45	1	0.87
OVARC1001161	64.57	91.5	118.13	1.42	1.83
OVARC1001162	67.07	66.3	60.17	0.99	0.9
OVARC1001163	44.06	41.7	37.81	0.95	0.91
OVARC1001167	131.97	178.27	124.95	1.35	0.95
OVARC1001169	36.78	27.43	38.99	1	1
OVARC1001170	65.61	77.2	57.31	1.18	0.87
OVARC1001171	222.01	381.42	266.16	1.72	1.2

(681)

特開 2002-191363

1359

1360

OVARC1001173	106.94	98.98	64.05	0.93	0.6
OVARC1001176	1738.1	1365.32	1450.41	0.79	0.83
OVARC1001180	170.76	219.17	160.37	1.28	0.94
OVARC1001188	43.78	31.39	45.77	0.91	1.05
OVARC1001200	30.93	28.46	25.56	1	1
OVARC1001202	89.8	109.06	76.14	1.21	0.85
OVARC1001206	25.21	19.96	14.96	1	1
OVARC1001209	135.99	167.39	86.92	1.23	0.64
OVARC1001219	43.42	67.27	69.27	1.55	1.6
OVARC1001222	51.14	44.76	64.42	0.88	1.26
OVARC1001232	87.58	109.93	82.19	1.26	0.94
OVARC1001240	62.57	67.43	59.2	1.08	0.95
OVARC1001243	31.53	23.79	40.34	1	1.01
OVARC1001244	71.67	92.27	74.16	1.29	1.03
OVARC1001246	183.36	317.23	328.1	1.73	1.79
OVARC1001247	38.53	38.34	42.68	1	1.07
OVARC1001260	40.46	31.35	41.53	0.99	1.03
OVARC1001261	29.3	27.69	37.42	1	1
OVARC1001268	156.38	139.16	131.87	0.89	0.84
OVARC1001270	29.69	25.77	25.2	1	1
OVARC1001271	61.41	90.42	61.05	1.47	0.99
OVARC1001282	21.59	10.83	20.59	1	1
OVARC1001296	23.87	32.76	62.54	1	1.56
OVARC1001306	17.53	22.01	16.97	1	1
OVARC1001314	28.21	26.99	26.93	1	1
OVARC1001316	34.31	39.56	30.84	1	1
OVARC1001329	232.27	293.46	231.75	1.26	1
OVARC1001330	21.47	21.46	24.72	1	1
OVARC1001336	68.81	83.92	59.46	1.22	0.86
OVARC1001338	23.84	21.6	27.85	1	1
OVARC1001339	166.35	209.33	201.15	1.26	1.21
OVARC1001340	24.69	30.23	15.34	1	1
OVARC1001341	83.51	53.41	58.61	0.64	0.7
OVARC1001342	906.51	940.44	1162.95	1.04	1.28
OVARC1001344	94.29	91.84	76.48	0.97	0.81
OVARC1001357	126.87	158.79	126.06	1.25	0.99
OVARC1001359	113.44	108.4	117.25	0.96	1.03
OVARC1001360	19.59	16.38	16.06	1	1
OVARC1001369	28.67	25.63	26.87	1	1
OVARC1001372	21.27	23.59	22.3	1	1
OVARC1001376	126.95	164.03	103.11	1.29	0.81
OVARC1001381	165.56	191.26	187.13	1.16	1.13
OVARC1001391	28.04	23.78	24.37	1	1
OVARC1001392	68.38	51.48	63.08	0.75	0.92
OVARC1001399	37.81	45.74	27	1.14	1
OVARC1001417	24.23	25.39	31.54	1	1
OVARC1001419	37.58	35.05	29.67	1	1
OVARC1001425	27.94	32.96	24.18	1	1
OVARC1001436	40.42	36.41	41.11	0.99	1.02
OVARC1001442	33.01	24.02	27.31	1	1

(682)

特開 2002-191363

1361

1362

OVARC1001451	84.3	90.12	71.71	1.07	0.85
OVARC1001452	38.24	38.53	36.34	1	1
OVARC1001453	38.85	54.09	45.75	1.35	1.14
OVARC1001476	60.15	81.57	84.23	1.36	1.4
OVARC1001480	26.66	26.16	25.2	1	1
OVARC1001489	48.15	49.45	101.95	1.03	2.12
OVARC1001493	41.75	42.93	31.69	1.03	0.96
OVARC1001496	60.58	93.19	72.72	1.54	1.2
OVARC1001499	73.29	81.63	65.54	1.11	0.89
OVARC1001506	64.89	70.58	57.55	1.09	0.89
OVARC1001509	55.61	77.79	48.65	1.4	0.87
OVARC1001510	26.95	24.97	15.31	1	1
OVARC1001516	38.07	44.29	47.82	1.11	1.2
OVARC1001525	23.96	35.03	30.42	1	1
OVARC1001542	67.08	99.57	88.99	1.48	1.33
OVARC1001544	95.76	116.73	98.74	1.22	1.03
OVARC1001546	49.52	43.47	38.3	0.88	0.81
OVARC1001547	23.53	18.95	25.34	1	1
OVARC1001555	1367.8	1096.38	1129.08	0.8	0.83
OVARC1001560	35.41	34.45	42.68	1	1.07
OVARC1001569	40.02	36.25	49.64	1	1.24
OVARC1001570	118.3	90.82	109.86	0.77	0.93
OVARC1001577	67.37	61.81	65.57	0.92	0.97
OVARC1001578	7.12	4.31	2.89	1	1
OVARC1001596	109.11	98.02	100.15	0.9	0.92
OVARC1001600	44.91	38.21	38.78	0.89	0.89
OVARC1001607	103.19	103.55	104.7	1	1.01
OVARC1001610	27.23	22.39	29.89	1	1
OVARC1001611	23.95	21.82	23.28	1	1
OVARC1001615	28.78	30.41	29.35	1	1
OVARC1001636	22.06	26.91	22.4	1	1
OVARC1001668	157.38	182.92	190.28	1.16	1.21
OVARC1001702	29.08	33.35	21.23	1	1
OVARC1001703	34.64	34.78	32.94	1	1
OVARC1001710	37	45.21	35.64	1.13	1
OVARC1001711	65.82	54.04	68.29	0.82	1.04
OVARC1001713	759.21	796.63	608.51	1.05	0.8
OVARC1001725	31.93	66.38	76.74	1.66	1.92
OVARC1001726	33.71	25.68	31.15	1	1
OVARC1001727	18.56	15.27	18.96	1	1
OVARC1001731	644.04	686.39	701.99	1.07	1.09
OVARC1001735	29.65	26.69	27.96	1	1
OVARC1001741	116.98	138.87	99.12	1.19	0.85
OVARC1001745	124.72	141.59	108.68	1.14	0.87
OVARC1001759	24.68	39.22	19.11	1	1
OVARC1001762	34.59	36.76	38.56	1	1
OVARC1001766	94.02	119.94	112.17	1.28	1.19
OVARC1001767	20.16	18.06	19.11	1	1
OVARC1001768	46.1	47.08	41.89	1.02	0.91
OVARC1001770	105.44	120.44	96.89	1.14	0.92

(683)

特開 2002-191363

1363

1364

OVARC1001776	23.68	21.65	15.61	1	1
OVARC1001791	28.43	27.18	17.64	1	1
OVARC1001795	19.53	18.07	13.49	1	1
OVARC1001798	101.84	142.85	110.21	1.4	1.08
OVARC1001802	65.06	78	62.46	1.2	0.96
OVARC1001805	35.52	48.94	64.35	1.22	1.61
OVARC1001807	56.61	69.76	47.03	1.23	0.83
OVARC1001809	1088.9	1547.3	919.46	1.42	0.84
OVARC1001812	39.57	54.09	36.17	1.35	1
OVARC1001813	58.4	60.12	50.34	1.03	0.86
OVARC1001820	46.93	54.05	53.52	1.15	1.14
OVARC1001828	22.82	20.17	17.42	1	1
OVARC1001833	23.13	23.75	22.25	1	1
OVARC1001839	25.87	24.28	22.99	1	1
OVARC1001846	32.88	39.67	35.78	1	1
OVARC1001849	42.93	45.6	36.21	1.06	0.93
OVARC1001861	38.57	44.42	39.11	1.11	1
OVARC1001873	21.53	26.72	31.29	1	1
OVARC1001879	31.66	30.09	34.2	1	1
OVARC1001880	41.78	56.41	43.72	1.35	1.05
OVARC1001883	54.76	53.84	51.98	0.98	0.95
OVARC1001900	41.08	39.22	37.45	0.97	0.97
OVARC1001901	32.69	29.55	28.05	1	1
OVARC1001911	24.17	21.88	19.26	1	1
OVARC1001916	40.38	47.57	41.31	1.18	1.02
OVARC1001928	23.54	15.33	21.01	1	1
OVARC1001937	116.75	121.57	137.57	1.04	1.18
OVARC1001940	27.84	23.38	26.49	1	1
OVARC1001942	51.05	48.07	54.43	0.94	1.07
OVARC1001943	55.91	66.85	90.22	1.2	1.61
OVARC1001949	49.03	55.03	45.81	1.12	0.93
OVARC1001950	111.23	119.03	88.58	1.07	0.8
OVARC1001952	925.52	837.31	491.71	0.9	0.53
OVARC1001954	22.54	22.58	19	1	1
OVARC1001963	39.61	39.13	33.68	1	1
OVARC1001983	156.69	178.79	170.9	1.14	1.09
OVARC1001987	47.79	48.7	50.85	1.02	1.06
OVARC1001989	115.86	105.75	98.03	0.91	0.85
OVARC1001991	52.33	46.51	46.57	0.89	0.89
OVARC1002005	125.8	138.41	116.88	1.1	0.93
OVARC1002044	114.35	98.85	116.74	0.86	1.02
OVARC1002046	159.32	196.2	173.16	1.23	1.09
OVARC1002050	40.99	40.28	45.96	0.98	1.12
OVARC1002058	37.98	49.57	48.1	1.24	1.2
OVARC1002066	144.74	132.59	118.22	0.92	0.82
OVARC1002082	144.14	154.11	143.8	1.07	1
OVARC1002091	52.45	54.17	45.92	1.03	0.88
OVARC1002092	32.76	33.97	32.14	1	1
OVARC1002093	66.9	77.69	43.57	1.16	0.65
OVARC1002094	25.97	23.94	18.11	1	1

(684)

特開 2002-191363

1365

1366

OVARC1002107	64.72	70.55	54.95	1.09	0.85
OVARC1002112	139.28	159.48	161.25	1.15	1.16
OVARC1002126	66.61	75.47	60.09	1.13	0.9
OVARC1002127	29.37	25.6	25.54	1	1
OVARC1002138	29.78	31.03	19.45	1	1
OVARC1002143	30.33	31.42	29.88	1	1
OVARC1002156	75.57	70.21	66.57	0.93	0.88
OVARC1002158	37.42	53.19	43.69	1.33	1.09
OVARC1002165	185.67	208.93	182.03	1.13	0.98
OVARC1002176	41.55	53.57	41.99	1.29	1.01
OVARC1002178	28.75	31.58	27.65	1	1
OVARC1002182	30.31	30.67	27.5	1	1
OVARC1002185	396.64	424.71	346.26	1.07	0.87
PLACE1000004	24.76	25.81	21.48	1	1
PLACE1000005	72.22	70.74	69.75	0.98	0.97
PLACE1000006	27.37	33.84	25.89	1	1
PLACE1000007	21.99	20.68	18.2	1	1
PLACE1000014	46.45	37.66	44.52	0.86	0.96
PLACE1000031	34.07	28.46	37.07	1	1
PLACE1000033	24.31	26.29	25.62	1	1
PLACE1000040	80.02	61.14	61.85	0.76	0.77
PLACE1000048	49.01	44.9	46.33	0.92	0.95
PLACE1000050	76.92	79.6	76.6	1.03	1
PLACE1000061	1833.7	1628.1	1862.77	0.89	1.02
PLACE1000066	100.49	137.52	143.02	1.37	1.42
PLACE1000075	27.62	28.66	37.82	1	1
PLACE1000078	108.32	94.57	102.84	0.87	0.95
PLACE1000081	22.97	26.54	22.74	1	1
PLACE1000086	55.62	56.11	45.41	1.01	0.82
PLACE1000094	22.42	45.59	26.18	1.14	1
PLACE1000101	70.88	63.96	84.2	0.9	1.19
PLACE1000121	24.82	20.96	22.84	1	1
PLACE1000133	96.36	136.13	146.32	1.41	1.52
PLACE1000142	38.87	34.64	40.54	1	1.01
PLACE1000146	23.88	32.9	31.49	1	1
PLACE1000163	55.99	61.61	42.54	1.1	0.76
PLACE1000172	34.86	30.15	28.35	1	1
PLACE1000181	28.93	39.59	30.35	1	1
PLACE1000184	31.13	34.98	39.62	1	1
PLACE1000185	59.19	53.44	60.92	0.9	1.03
PLACE1000198	30.12	40.19	26.16	1	1
PLACE1000213	38.99	48.17	37.26	1.2	1
PLACE1000214	34.7	46.78	49.53	1.17	1.24
PLACE1000220	73.76	70.27	66.16	0.95	0.9
PLACE1000231	95.59	80.27	84.2	0.84	0.88
PLACE1000236	39.98	42.02	37.68	1.05	1
PLACE1000245	82.41	100.11	79.89	1.21	0.97
PLACE1000246	40.74	34.37	31.89	0.98	0.98
PLACE1000258	130.7	150.85	139.27	1.15	1.07
PLACE1000288	24.77	24.09	32.17	1	1

(685)

特開 2002-191363

1367

1368

PLACE1000292	106.53	113.38	98.12	1.06	0.92
PLACE1000302	18.27	27.21	15.42	1	1
PLACE1000304	65.7	62.07	62.7	0.94	0.95
PLACE1000308	32.53	34.89	34.8	1	1
PLACE1000309	58.44	64.89	69.33	1.11	1.19
PLACE1000312	26.22	23.84	24.8	1	1
PLACE1000330	24.35	26.43	20.16	1	1
PLACE1000332	35.75	32.57	26.57	1	1
PLACE1000347	42.57	48.19	57.82	1.13	1.36
PLACE1000351	31.19	20.96	22.08	1	1
PLACE1000374	85.05	103.62	71.9	1.22	0.85
PLACE1000380	36.69	40.89	43.75	1.02	1.09
PLACE1000383	27.18	37.69	30.62	1	1
PLACE1000397	29.25	23.24	30.83	1	1
PLACE1000401	38.53	42.86	33.59	1.07	1
PLACE1000406	33.82	36.73	38.14	1	1
PLACE1000412	36.43	38.07	40.64	1	1.02
PLACE1000420	89.72	84	76.38	0.94	0.85
PLACE1000421	48.98	53.73	40.47	1.1	0.83
PLACE1000423	289.62	300.74	318.77	1.04	1.1
PLACE1000424	47.59	56.91	37.74	1.2	0.84
PLACE1000430	35.21	29.9	25.42	1	1
PLACE1000433	31.4	33.71	27.27	1	1
PLACE1000435	67.53	71.28	61.52	1.06	0.91
PLACE1000437	85.98	76.05	90.26	0.88	1.05
PLACE1000442	71.69	83.76	65.33	1.17	0.91
PLACE1000444	281.96	322.3	237.42	1.14	0.84
PLACE1000453	93.48	96.07	86.64	1.03	0.93
PLACE1000456	155.96	197.31	251.5	1.27	1.61
PLACE1000465	88.14	80.44	56.34	0.91	0.64
PLACE1000481	60.69	64.48	61.62	1.06	1.02
PLACE1000492	29.41	31.26	28.26	1	1
PLACE1000508	43.75	36.51	33.48	0.91	0.91
PLACE1000512	43.13	37.22	29.76	0.93	0.93
PLACE1000540	60.89	54.69	53.42	0.9	0.88
PLACE1000541	194.23	186.38	150.17	0.96	0.77
PLACE1000546	58.5	58.41	50.88	1	0.87
PLACE1000547	62.12	74.88	58.14	1.21	0.94
PLACE1000560	27.65	32.31	26.9	1	1
PLACE1000562	68.1	102.88	93.56	1.51	1.37
PLACE1000564	32.94	44.95	38.66	1.12	1
PLACE1000583	129.86	146.67	93.02	1.13	0.72
PLACE1000587	76.59	77.27	62.7	1.01	0.82
PLACE1000588	44.17	50.72	31.24	1.15	0.91
PLACE1000596	64.54	70.96	67.19	1.1	1.04
PLACE1000599	121.52	140.66	95.47	1.16	0.79
PLACE1000605	111.2	111.23	135.21	1	1.22
PLACE1000610	36.84	46.4	33.01	1.16	1
PLACE1000611	88.85	115.78	96.74	1.3	1.09
PLACE1000626	81.31	97.7	63.54	1.2	0.78

(686)

特開 2002-191363

1369

1370

PLACE1000633	89.73	103.53	69	1.15	0.77
PLACE1000636	43.66	38.64	31.59	0.92	0.92
PLACE1000653	48.6	59.21	37.82	1.22	0.82
PLACE1000656	76.78	115.05	89.87	1.5	1.17
PLACE1000663	22.81	25.89	22.83	1	1
PLACE1000706	47.62	68.55	44.03	1.44	0.92
PLACE1000712	42.96	41.85	44.5	0.97	1.04
PLACE1000716	36.32	30.57	37.12	1	1
PLACE1000740	74.99	71.02	72.48	0.95	0.97
PLACE1000748	59.7	44.52	53.83	0.75	0.9
PLACE1000749	69.67	71.54	88.13	1.03	1.26
PLACE1000751	90.04	63.66	96.59	0.71	1.07
PLACE1000755	28.06	18.71	33.71	1	1
PLACE1000769	28.38	22.38	36.57	1	1
PLACE1000778	309.15	302.88	359.07	0.98	1.16
PLACE1000785	281.33	264.97	226.92	0.94	0.81
PLACE1000786	44.9	51.88	43.46	1.16	0.97
PLACE1000793	49.42	67.76	62.97	1.37	1.27
PLACE1000795	44.34	51.39	45.61	1.16	1.03
PLACE1000798	56	57.48	53.18	1.03	0.95
PLACE1000812	50.83	52.85	42.72	1.04	0.84
PLACE1000823	67.58	81.4	78.53	1.2	1.16
PLACE1000825	292.82	278.76	227.12	0.95	0.78
PLACE1000838	117.7	102.23	110.08	0.87	0.94
PLACE1000841	40.32	63.69	47.74	1.58	1.18
PLACE1000843	46.51	40.25	48.16	0.87	1.04
PLACE1000849	46.45	41.04	46.47	0.88	1
PLACE1000856	36.71	40.33	41.25	1.01	1.03
PLACE1000863	62.44	65.02	86.88	1.04	1.39
PLACE1000876	51.73	69.32	56.48	1.34	1.09
PLACE1000899	40.08	45.46	36.02	1.13	1
PLACE1000907	121.51	131.23	137.97	1.08	1.14
PLACE1000909	64.51	62.85	76.14	0.97	1.18
PLACE1000912	17.7	22.05	17.88	1	1
PLACE1000914	32.84	31.69	26.59	1	1
PLACE1000918	23.6	42.91	25.79	1.07	1
PLACE1000927	92.92	134.99	144.28	1.45	1.55
PLACE1000931	43.54	48.87	49.05	1.12	1.13
PLACE1000944	46.12	56.25	42.79	1.22	0.93
PLACE1000948	31.87	30.6	29.54	1	1
PLACE1000958	26.58	22.19	27.28	1	1
PLACE1000972	77.78	102.18	78.45	1.31	1.01
PLACE1000977	37.49	43.04	37.27	1.08	1
PLACE1000979	56.17	42.68	63.84	0.76	1.14
PLACE1000986	25.37	30.46	29.36	1	1
PLACE1000987	31.24	34.96	33.17	1	1
PLACE1001000	60.23	87.53	55.33	1.45	0.92
PLACE1001007	62.89	63.91	57.94	1.02	0.92
PLACE1001010	32.73	43.47	36.9	1.09	1
PLACE1001015	25.94	37.23	23.1	1	1

(687)

特開 2002-191363

1371

1372

PLACE1001016	67.56	68.13	64.74	1.01	0.96
PLACE1001022	24.1	29.77	22.61	1	1
PLACE1001024	24.54	28.89	23.58	1	1
PLACE1001036	52.29	53.52	38.48	1.02	0.76
PLACE1001038	253.42	281.94	250.92	1.11	0.99
PLACE1001048	26.43	26.89	24.61	1	1
PLACE1001054	1175.5	1243.17	853.43	1.06	0.73
PLACE1001062	65.69	88.71	61.11	1.35	0.93
PLACE1001063	22.22	18.82	21.06	1	1
PLACE1001076	23.43	22.79	18.47	1	1
PLACE1001081	72.77	65.53	103.11	0.9	1.42
PLACE1001088	32.05	30.28	39.19	1	1
PLACE1001092	53.83	55.44	46.76	1.03	0.87
PLACE1001098	151.65	175.93	181.39	1.16	1.2
PLACE1001100	54.47	53.59	38.53	0.98	0.73
PLACE1001104	32.63	33.73	33.87	1	1
PLACE1001114	46.98	47.37	50.39	1.01	1.07
PLACE1001118	53.9	66.84	51.97	1.24	0.96
PLACE1001123	66.86	69.57	68.68	1.04	1.03
PLACE1001136	65.85	79.52	60.64	1.21	0.92
PLACE1001144	90.6	79.44	58.29	0.88	0.64
PLACE1001147	55.83	58.23	55.99	1.04	1
PLACE1001148	41.47	34.75	55.19	0.96	1.33
PLACE1001159	31.15	31.69	25.91	1	1
PLACE1001168	41.99	53.56	68.74	1.28	1.64
PLACE1001171	18.09	17.01	11.3	1	1
PLACE1001183	25.6	35.65	26.27	1	1
PLACE1001185	74.4	83.96	71.65	1.13	0.96
PLACE1001201	65.89	45.96	32.25	0.7	0.61
PLACE1001229	79.77	73.37	75.05	0.92	0.94
PLACE1001231	49.09	70.09	47.97	1.43	0.98
PLACE1001238	59	59.58	44.13	1.01	0.75
PLACE1001241	37.8	28.25	39	1	1
PLACE1001242	212.52	280.16	306.05	1.32	1.44
PLACE1001247	39.43	48.02	40.05	1.2	1
PLACE1001250	43.9	48.93	49.67	1.11	1.13
PLACE1001257	69.88	87.14	49.9	1.25	0.71
PLACE1001272	62.08	59.2	49.01	0.95	0.79
PLACE1001279	36.9	28.46	28.45	1	1
PLACE1001280	59.82	68.95	57.17	1.15	0.96
PLACE1001294	55.4	76.7	58.04	1.38	1.05
PLACE1001295	37.73	42.1	34.43	1.05	1
PLACE1001300	38.62	81.94	59.07	2.05	1.48
PLACE1001304	76.32	107.22	96.84	1.4	1.27
PLACE1001311	71.6	81.41	59.85	1.14	0.84
PLACE1001323	72.17	76.01	59.14	1.05	0.82
PLACE1001325	59.21	89.16	52.84	1.51	0.89
PLACE1001340	59.98	70.04	68.95	1.17	1.15
PLACE1001344	16.91	17.79	16.63	1	1
PLACE1001351	37.9	35.39	36.71	1	1

(688)

特開 2002-191363

1373

1374

PLACE1001366	39.46	53.03	43.75	1.33	1.09
PLACE1001377	39.42	36.27	35.5	1	1
PLACE1001383	50.96	69.88	43.67	1.37	0.86
PLACE1001384	29.93	24.61	29.44	1	1
PLACE1001387	19.84	23.39	22.61	1	1
PLACE1001395	48.1	54.54	59.91	1.13	1.25
PLACE1001399	73	89.28	87.29	1.22	1.2
PLACE1001401	17.36	10.16	14.75	1	1
PLACE1001407	173.32	153.48	184.52	0.89	1.06
PLACE1001412	28.36	51.42	50.36	1.29	1.26
PLACE1001414	103.73	119.31	104.84	1.15	1.01
PLACE1001416	45.34	37.52	46.55	0.88	1.03
PLACE1001433	686.08	633.61	710.08	0.92	1.03
PLACE1001440	31.5	30.51	37.83	1	1
PLACE1001456	93.44	86.92	92.14	0.93	0.99
PLACE1001464	144.35	156.63	222.58	1.09	1.54
PLACE1001468	21.17	20.97	23.23	1	1
PLACE1001484	39.26	44.3	44.46	1.11	1.11
PLACE1001500	29.64	27.34	25.08	1	1
PLACE1001502	44.35	52.36	37.68	1.18	0.9
PLACE1001503	106.67	108.23	94.51	1.01	0.89
PLACE1001505	107.02	84.34	109.11	0.79	1.02
PLACE1001513	59.63	60.93	69.28	1.02	1.16
PLACE1001516	46.89	44.32	37.19	0.95	0.85
PLACE1001517	67.5	61.74	72.29	0.91	1.07
PLACE1001523	35.36	145.54	176.62	3.64	4.42
PLACE1001526	67.64	68.66	60.16	1.02	0.89
PLACE1001534	59.24	49.05	59.78	0.83	1.01
PLACE1001536	21.36	23	24.62	1	1
PLACE1001545	299.5	242.39	336.18	0.81	1.12
PLACE1001551	52.49	59.71	48.37	1.14	0.92
PLACE1001564	32.98	43.9	34.74	1.1	1
PLACE1001570	37.67	45.32	43.97	1.13	1.1
PLACE1001571	60.72	58.01	57.65	0.96	0.95
PLACE1001595	131.76	206.77	116.03	1.57	0.88
PLACE1001602	140.17	124.26	134.66	0.89	0.96
PLACE1001603	106.03	99.78	98.98	0.94	0.93
PLACE1001608	50.87	50.16	41.39	0.99	0.81
PLACE1001610	152.25	167.91	138.81	1.1	0.91
PLACE1001611	23.89	32.27	24.71	1	1
PLACE1001629	26.65	29.58	37.06	1	1
PLACE1001632	30.66	37.71	44.36	1	1.11
PLACE1001634	172.71	157.55	99.09	0.91	0.57
PLACE1001637	25.7	21.53	18.93	1	1
PLACE1001640	57.73	81.27	57.98	1.41	1
PLACE1001655	20.38	17.61	16.27	1	1
PLACE1001672	36.77	44.13	45.88	1.1	1.15
PLACE1001676	61.14	47.49	68.28	0.78	1.12
PLACE1001683	130.38	163.09	126.67	1.25	0.97
PLACE1001691	66.24	137.29	148.77	2.07	2.25

(689)

特開 2002-191363

1375

1376

PLACE1001692	107.1	130.47	90.19	1.22	0.84
PLACE1001705	103	105.83	98.71	1.03	0.96
PLACE1001716	34.04	37.84	33.36	1	1
PLACE1001720	22.82	19.21	23.04	1	1
PLACE1001728	23.74	33.98	26.51	1	1
PLACE1001729	42.25	41.32	31.82	0.98	0.95
PLACE1001739	27.27	36.66	35.51	1	1
PLACE1001740	41.28	37.48	41.21	0.97	1
PLACE1001745	33.2	30.85	25.22	1	1
PLACE1001746	58.49	62.26	51.19	1.06	0.88
PLACE1001748	91.44	111.27	90.04	1.22	0.98
PLACE1001753	46.48	41.49	42.81	0.89	0.92
PLACE1001756	206.24	206.45	146.37	1	0.71
PLACE1001761	105	105.33	114.94	1	1.09
PLACE1001767	906.39	1018.83	646.39	1.12	0.71
PLACE1001771	50.78	47.05	50.31	0.93	0.99
PLACE1001775	56.72	58.09	52.35	1.02	0.92
PLACE1001777	250.62	301.05	385.02	1.2	1.54
PLACE1001781	47.54	45.66	50.36	0.96	1.06
PLACE1001783	25.79	23.91	24.74	1	1
PLACE1001786	27.32	22.95	23.3	1	1
PLACE1001788	63.8	50.58	49.3	0.79	0.77
PLACE1001795	55.25	47.94	42.33	0.87	0.77
PLACE1001799	19.31	21.26	19.78	1	1
PLACE1001810	37.78	34	28.42	1	1
PLACE1001817	51.28	35.63	49.47	0.78	0.96
PLACE1001821	90.72	88.49	68.61	0.98	0.76
PLACE1001836	40.93	47.62	45.4	1.16	1.11
PLACE1001844	50.7	52.62	49.15	1.04	0.97
PLACE1001845	114.64	148.46	138.06	1.3	1.2
PLACE1001858	39.69	37.7	45.91	1	1.15
PLACE1001869	66.67	56.7	50.32	0.85	0.75
PLACE1001890	139.54	181.8	133.9	1.3	0.96
PLACE1001897	103.15	90.08	92.86	0.87	0.9
PLACE1001902	132.93	111.17	101.84	0.84	0.77
PLACE1001904	21.63	26.18	17.37	1	1
PLACE1001907	51.16	54.84	43.55	1.07	0.85
PLACE1001910	4125.5	5812.98	5956.63	1.41	1.44
PLACE1001912	98.39	125.69	81.71	1.28	0.83
PLACE1001918	75.3	178.35	171.51	2.37	2.28
PLACE1001920	38.58	44.62	25.77	1.12	1
PLACE1001928	59.67	54.32	45.81	0.91	0.77
PLACE1001930	42.5	39.89	29.86	0.94	0.94
PLACE1001949	30.76	31.17	25.06	1	1
PLACE1001959	30.92	27.74	44.02	1	1.1
PLACE1001969	43.67	68.4	60.61	1.57	1.39
PLACE1001974	58.47	87.97	70.98	1.5	1.21
PLACE1001981	20.55	19.1	19.17	1	1
PLACE1001983	50.42	56.62	39.13	1.12	0.79
PLACE1001989	64.23	61.57	57.16	0.96	0.89

(690)

特開 2002-191363

1377

1378

PLACE1002004	138.52	136.78	156.35	0.99	1.13
PLACE1002008	78.06	75.77	80.18	0.97	1.03
PLACE1002015	397.49	417.71	445.16	1.05	1.12
PLACE1002044	41.16	31.51	40.22	0.97	0.98
PLACE1002046	44.29	35.77	47.55	0.9	1.07
PLACE1002052	20.68	23.54	25.31	1	1
PLACE1002066	178.37	183.05	183.25	1.03	1.03
PLACE1002072	78.28	81.5	77.13	1.04	0.99
PLACE1002073	27.9	28.71	34.29	1	1
PLACE1002080	61.82	72.61	63.14	1.17	1.02
PLACE1002081	40.52	38.51	38.8	0.99	0.99
PLACE1002090	79.31	102.85	131.29	1.3	1.66
PLACE1002095	63.25	73	81.15	1.15	1.28
PLACE1002102	47.9	64.22	87.09	1.34	1.82
PLACE1002109	121.13	154.2	113.37	1.27	0.94
PLACE1002115	47.16	53.43	44.36	1.13	0.94
PLACE1002119	220.58	222	254.75	1.01	1.15
PLACE1002140	224.21	154.92	220.12	0.69	0.98
PLACE1002150	46.55	38.95	33.85	0.86	0.86
PLACE1002153	34.37	46.95	43.15	1.17	1.08
PLACE1002157	36.71	37.17	35.5	1	1
PLACE1002163	41.57	43.98	53.8	1.06	1.29
PLACE1002168	80.08	72.26	77.83	0.9	0.97
PLACE1002170	32.62	28.6	31.69	1	1
PLACE1002171	54.23	42.59	50.12	0.79	0.92
PLACE1002180	55.02	46.62	41.61	0.85	0.76
PLACE1002184	117.06	172.52	143.25	1.47	1.22
PLACE1002200	33.2	36.06	34.77	1	1
PLACE1002205	93.98	75.49	65.05	0.8	0.69
PLACE1002213	49.41	60.19	55.06	1.22	1.11
PLACE1002219	32.08	34.58	42.83	1	1.07
PLACE1002227	39.22	41.44	39.22	1.04	1
PLACE1002253	13.18	15.01	11.64	1	1
PLACE1002256	36.08	42.07	41.31	1.05	1.03
PLACE1002259	35.42	38.71	28.99	1	1
PLACE1002285	21.44	22.04	22.02	1	1
PLACE1002301	39.97	51.82	47.69	1.3	1.19
PLACE1002310	38.3	53.71	59.83	1.34	1.5
PLACE1002311	32.31	31.35	21.52	1	1
PLACE1002319	39.1	37.88	38.01	1	1
PLACE1002329	42.35	45.03	28.89	1.06	0.94
PLACE1002333	22.15	21.3	18.15	1	1
PLACE1002342	57.91	78.12	77.69	1.35	1.34
PLACE1002343	24.89	27.37	21.38	1	1
PLACE1002355	29.14	51.73	37.92	1.29	1
PLACE1002358	35.63	34.51	30.06	1	1
PLACE1002359	59.86	66.1	56.59	1.1	0.95
PLACE1002374	108.32	173.07	169.76	1.6	1.57
PLACE1002376	123.21	156.84	143.37	1.27	1.16
PLACE1002379	56.01	73.79	77.99	1.32	1.39

(691)

特開 2002-191363

1379

1380

PLACE1002386	18.78	17.51	20.81	1	1
PLACE1002395	131.23	134.81	90.74	1.03	0.69
PLACE1002399	33.82	41.2	50.11	1.03	1.25
PLACE1002407	28.08	24.72	32.3	1	1
PLACE1002433	39.73	34.06	26.39	1	1
PLACE1002437	28.87	27.38	24.22	1	1
PLACE1002438	27.94	29.9	24.65	1	1
PLACE1002446	31.05	33.92	52.9	1	1.32
PLACE1002447	38.57	23.64	33.34	1	1
PLACE1002450	31.98	40.64	39.26	1.02	1
PLACE1002462	34	36.24	27.07	1	1
PLACE1002465	37.59	32.41	32.62	1	1
PLACE1002474	34.19	30.1	31.83	1	1
PLACE1002477	215.53	256.98	192.24	1.19	0.89
PLACE1002493	117.08	120.94	66.9	1.03	0.57
PLACE1002497	22.96	20.27	15.65	1	1
PLACE1002499	50.73	58.7	59.23	1.16	1.17
PLACE1002500	96.27	147.8	112.4	1.54	1.17
PLACE1002514	33.61	26.45	23.6	1	1
PLACE1002518	41.61	56.08	61.21	1.35	1.47
PLACE1002529	35.95	30.55	27.67	1	1
PLACE1002532	111.18	127.2	70.27	1.14	0.63
PLACE1002536	154.38	153.55	128.51	0.99	0.83
PLACE1002537	61.55	59.88	57.19	0.97	0.93
PLACE1002539	37.9	30.66	31.19	1	1
PLACE1002547	67.73	93.86	89.6	1.39	1.32
PLACE1002571	63.4	74.13	58.68	1.17	0.93
PLACE1002578	132.43	150.29	116.55	1.13	0.88
PLACE1002583	32.14	34.49	24.56	1	1
PLACE1002591	37.57	41	27.91	1.03	1
PLACE1002598	96.18	87.39	79.55	0.91	0.83
PLACE1002604	54.28	66.18	49.22	1.22	0.91
PLACE1002612	95.35	95.19	103.19	1	1.08
PLACE1002625	35.29	35.5	37.17	1	1
PLACE1002638	36.08	51.42	37.95	1.29	1
PLACE1002655	77.69	90.45	87.49	1.16	1.13
PLACE1002665	31.49	32.82	23.4	1	1
PLACE1002685	17.89	18.59	13.37	1	1
PLACE1002692	127.12	135.43	105.27	1.07	0.83
PLACE1002714	19.26	20.24	19.68	1	1
PLACE1002721	34.62	40.72	46.88	1.02	1.17
PLACE1002722	19.06	20.66	14.68	1	1
PLACE1002726	40.31	30.8	31.87	0.99	0.99
PLACE1002756	78.21	83.19	71.78	1.06	0.92
PLACE1002768	25.22	31.86	26.26	1	1
PLACE1002772	20.72	18	15.78	1	1
PLACE1002775	68.56	83.59	84.85	1.22	1.24
PLACE1002780	19.13	20.21	19.82	1	1
PLACE1002782	14.99	11.76	15.12	1	1
PLACE1002794	26.71	32.3	25.11	1	1

(692)

特開 2002-191363

1381

1382

PLACE1002795	51.53	51.44	55.42	1	1.08
PLACE1002811	19.7	15.5	14.68	1	1
PLACE1002815	51.13	46.51	44.21	0.91	0.86
PLACE1002816	144.66	145.59	134.13	1.01	0.93
PLACE1002822	37.08	39.75	33.18	1	1
PLACE1002833	99.65	99.15	105.52	0.99	1.06
PLACE1002834	90.65	105.86	92.19	1.17	1.02
PLACE1002835	21.6	20.73	17.21	1	1
PLACE1002839	35.83	40.12	42.4	1	1.06
PLACE1002851	37.62	50.62	39.26	1.27	1
PLACE1002853	77.21	86.13	73.13	1.12	0.95
PLACE1002881	178.57	217.28	143.56	1.22	0.8
PLACE1002901	162.03	130.28	155.87	0.8	0.96
PLACE1002904	41.21	32.91	35.47	0.97	0.97
PLACE1002905	60.55	65.83	58.21	1.09	0.96
PLACE1002908	37.5	41.43	40.54	1.04	1.01
PLACE1002911	50.17	67.78	80.71	1.35	1.61
PLACE1002941	41.63	57.62	41.65	1.38	1
PLACE1002950	60.18	35.74	54.95	0.66	0.91
PLACE1002955	500.04	509.35	572.07	1.02	1.14
PLACE1002958	300.19	264.52	323.01	0.88	1.08
PLACE1002962	31.57	28.46	29.19	1	1
PLACE1002967	58.45	68.94	55.81	1.18	0.95
PLACE1002968	74.85	90.97	72.09	1.22	0.96
PLACE1002976	111.45	136.26	171.19	1.22	1.54
PLACE1002991	68.54	89.4	62.41	1.3	0.91
PLACE1002993	109.46	111.94	93.86	1.02	0.86
PLACE1002996	52.75	45.18	57.04	0.86	1.08
PLACE1003010	227.78	176.08	214.3	0.77	0.94
PLACE1003025	82.24	84.89	67.49	1.03	0.82
PLACE1003027	43.36	36.79	58.67	0.92	1.35
PLACE1003044	27.11	31.6	30.76	1	1
PLACE1003045	28.39	20.72	24.75	1	1
PLACE1003052	46.7	55.42	35.28	1.19	0.86
PLACE1003083	39.42	35.4	30.63	1	1
PLACE1003085	105.31	95.13	165.83	0.9	1.57
PLACE1003092	53.65	49.63	61.79	0.93	1.15
PLACE1003097	19.84	17.97	16.61	1	1
PLACE1003100	43.48	42.63	54.49	0.98	1.25
PLACE1003108	67.4	69.86	56.14	1.04	0.83
PLACE1003115	2429.8	1938.41	1364	0.8	0.56
PLACE1003120	190.38	186.69	172.7	0.98	0.91
PLACE1003135	22.74	21.24	19.25	1	1
PLACE1003136	66.07	82.56	84.67	1.25	1.28
PLACE1003141	38.76	48.18	41.93	1.2	1.05
PLACE1003145	39.3	41.02	42.47	1.03	1.06
PLACE1003147	32.84	46.21	44.01	1.16	1.1
PLACE1003153	55.32	63.18	48.47	1.14	0.88
PLACE1003163	87.96	126.75	137.45	1.44	1.56
PLACE1003172	292.86	334.87	290.66	1.14	0.99

(693)

特開 2002-191363

1383

1384

PLACE1003174	44.47	47.73	42.71	1.07	0.96
PLACE1003176	35.28	43.34	27.58	1.08	1
PLACE1003181	44.5	38.93	34.82	0.9	0.9
PLACE1003184	21.75	19.42	16.07	1	1
PLACE1003190	53.54	44.24	78.5	0.83	1.47
PLACE1003200	19.44	17.3	16.03	1	1
PLACE1003205	149.26	188.41	143.32	1.26	0.96
PLACE1003209	24.57	23.64	19.44	1	1
PLACE1003214	29.44	37.32	25.25	1	1
PLACE1003229	58.97	74.77	48.92	1.27	0.83
PLACE1003238	23.79	31.2	22.95	1	1
PLACE1003249	76.24	88.55	60.02	1.16	0.79
PLACE1003256	136.88	173.8	158.56	1.27	1.16
PLACE1003258	26.94	29.08	19.67	1	1
PLACE1003279	289.62	288.28	218.59	1	0.75
PLACE1003294	40.84	31.78	29.14	0.98	0.98
PLACE1003296	50.97	44.86	42.34	0.88	0.83
PLACE1003297	368.37	299.91	291.38	0.81	0.79
PLACE1003302	113.04	132.54	114.02	1.17	1.01
PLACE1003334	41.77	44.78	39.07	1.07	0.96
PLACE1003337	313.35	296.56	240.71	0.95	0.77
PLACE1003342	22.4	35.38	23.63	1	1
PLACE1003343	30.97	30.24	26.77	1	1
PLACE1003344	199.37	235.82	176.95	1.18	0.89
PLACE1003353	498.58	500.44	453.52	1	0.91
PLACE1003361	93.89	106.31	91.43	1.13	0.97
PLACE1003366	78.26	94.5	66.92	1.21	0.86
PLACE1003369	5.77	2.72	5.34	1	1
PLACE1003372	57.21	51.02	82.26	0.89	1.44
PLACE1003373	193.53	234.31	178.93	1.21	0.92
PLACE1003375	31.83	33.45	28.49	1	1
PLACE1003378	14.4	11.05	6.95	1	1
PLACE1003383	35.89	36.91	29.46	1	1
PLACE1003394	68.37	65.46	64.47	0.96	0.94
PLACE1003401	9.48	14.44	5.74	1	1
PLACE1003405	31.77	38.58	39.98	1	1
PLACE1003407	47.62	69.36	59.4	1.46	1.25
PLACE1003420	97.91	112.98	116.4	1.15	1.19
PLACE1003428	14.9	21.6	14.87	1	1
PLACE1003432	54.93	40.52	47.7	0.74	0.87
PLACE1003438	32.24	30.44	24.69	1	1
PLACE1003452	46.53	35.85	47.65	0.86	1.02
PLACE1003454	39.61	50.18	45.74	1.25	1.14
PLACE1003455	36.33	33.93	32.61	1	1
PLACE1003456	108.71	98.84	105.2	0.91	0.97
PLACE1003460	102.86	100.68	102.3	0.98	0.99
PLACE1003478	21.22	36.98	21.27	1	1
PLACE1003484	350.57	284.4	235.73	0.81	0.67
PLACE1003493	206.05	248.77	248.38	1.21	1.21
PLACE1003503	782.31	770.43	847.28	0.98	1.08

(694)

特開 2002-191363

1385

1386

PLACE1003505	183.24	231.77	199.38	1.26	1.09
PLACE1003516	45.11	53.26	56.23	1.18	1.25
PLACE1003519	314.62	383.46	434.26	1.22	1.38
PLACE1003520	262.65	580.45	525.45	2.21	2
PLACE1003521	49.86	76.87	55.98	1.54	1.12
PLACE1003525	125.96	134.81	112.91	1.07	0.9
PLACE1003528	1564.6	1648.77	1526.57	1.05	0.98
PLACE1003529	41.65	28.28	28.22	0.96	0.96
PLACE1003537	58.26	41.57	70.53	0.71	1.21
PLACE1003549	71.19	60.94	46.62	0.86	0.65
PLACE1003553	33.32	33.26	30.39	1	1
PLACE1003566	129.04	183.13	165.12	1.42	1.28
PLACE1003568	25.89	43.04	31.17	1.08	1
PLACE1003573	50.28	42.88	45.95	0.85	0.91
PLACE1003575	122.7	135.29	90.48	1.1	0.74
PLACE1003583	35.13	38.24	32.11	1	1
PLACE1003584	68.08	74.06	61.63	1.09	0.91
PLACE1003592	130.19	171.31	138.87	1.32	1.07
PLACE1003593	31.68	37.2	39.66	1	1
PLACE1003594	71.43	93.99	104.68	1.32	1.47
PLACE1003596	150.3	154.27	135.36	1.03	0.9
PLACE1003598	100.52	98.24	96.34	0.98	0.96
PLACE1003602	52.87	50.33	60.7	0.95	1.15
PLACE1003605	467.51	490.79	396.84	1.05	0.85
PLACE1003611	89.33	110.21	81.69	1.23	0.91
PLACE1003618	28.29	31.43	27.78	1	1
PLACE1003625	31.69	39.67	45.94	1	1.15
PLACE1003626	283.72	224.78	209.61	0.79	0.74
PLACE1003630	54.08	63.75	40.92	1.18	0.76
PLACE1003635	39.71	34.86	29.62	1	1
PLACE1003638	65.67	77.09	51.77	1.17	0.79
PLACE1003644	148.25	167.05	116.51	1.13	0.79
PLACE1003654	29.17	27.24	37.13	1	1
PLACE1003656	45.98	64.88	74.77	1.41	1.63
PLACE1003660	69.53	86.73	71.97	1.25	1.04
PLACE1003669	32.51	45.27	35.34	1.13	1
PLACE1003670	155.29	157.75	143.9	1.02	0.93
PLACE1003671	29.58	31.07	29.68	1	1
PLACE1003697	32.77	42.06	124.21	1.05	3.11
PLACE1003704	46.2	53.43	49.07	1.16	1.06
PLACE1003709	21.1	22.79	22.2	1	1
PLACE1003711	57.95	44.93	49.55	0.78	0.86
PLACE1003723	64.95	63.87	54.85	0.98	0.84
PLACE1003724	126.19	128.74	117.29	1.02	0.93
PLACE1003737	33.88	40.76	32.52	1.02	1
PLACE1003738	56.01	62.95	50.36	1.12	0.9
PLACE1003742	59.61	66.3	57.81	1.11	0.97
PLACE1003744	87.46	98.48	85.09	1.13	0.97
PLACE1003758	26.25	23.92	30.05	1	1
PLACE1003760	198.63	264.05	284.47	1.33	1.43

(695)

特開 2002-191363

1387

1388

PLACE1003762	48.59	59.1	41.22	1.22	0.85
PLACE1003765	72.13	74.99	56.69	1.04	0.79
PLACE1003768	52.93	50.85	38.01	0.96	0.76
PLACE1003771	156.89	168.16	133.56	1.07	0.85
PLACE1003772	1326.1	1338.01	1235.63	1.01	0.93
PLACE1003783	40.78	42.6	38.87	1.04	0.98
PLACE1003784	35.32	31.33	31.62	1	1
PLACE1003788	16.15	15.6	13.57	1	1
PLACE1003795	69.33	75.16	71.21	1.08	1.03
PLACE1003827	74.21	106.07	131.59	1.43	1.77
PLACE1003833	40.41	38.8	30.6	0.99	0.99
PLACE1003839	564.98	498.2	452.27	0.88	0.8
PLACE1003845	107.06	114.29	87.76	1.07	0.82
PLACE1003850	63.8	58.95	53.45	0.92	0.84
PLACE1003852	24.02	15.1	18.58	1	1
PLACE1003858	29.1	40.09	42.99	1	1.07
PLACE1003861	112.21	124.5	120.75	1.11	1.08
PLACE1003864	40.19	41.75	36.75	1.04	1
PLACE1003870	266.48	329.04	207.77	1.23	0.78
PLACE1003885	25.52	42.25	33.9	1.06	1
PLACE1003886	81.35	79.28	74.9	0.97	0.92
PLACE1003888	39.04	47.84	30.42	1.2	1
PLACE1003892	206.13	289.84	210.39	1.41	1.02
PLACE1003900	57.88	76.01	53.39	1.31	0.92
PLACE1003902	40.78	41.51	43.92	1.02	1.08
PLACE1003903	28.5	35.54	27.43	1	1
PLACE1003915	40.52	47.1	35.74	1.16	0.99
PLACE1003918	52.38	76.45	47.24	1.46	0.9
PLACE1003923	53.3	58.03	49.05	1.09	0.92
PLACE1003932	49.92	74.44	58.48	1.49	1.17
PLACE1003936	29.36	44.28	34.72	1.11	1
PLACE1003966	46.44	61.08	55.25	1.32	1.19
PLACE1003968	62.13	67.95	76.19	1.09	1.23
PLACE1004018	65.03	72.67	69.12	1.12	1.06
PLACE1004020	100.08	107.43	83.05	1.07	0.83
PLACE1004028	14.25	14.56	14.5	1	1
PLACE1004034	37.37	40.65	43.69	1.02	1.09
PLACE1004042	347.79	322.05	361.95	0.93	1.04
PLACE1004078	43.15	47.81	51.52	1.11	1.19
PLACE1004103	74.8	77.55	95.57	1.04	1.28
PLACE1004104	204.1	203.04	177.57	0.99	0.87
PLACE1004113	51.84	54.36	57.12	1.05	1.1
PLACE1004114	73.67	91.18	74.31	1.24	1.01
PLACE1004118	28.97	31.53	31.34	1	1
PLACE1004128	64.64	48.58	71.02	0.75	1.1
PLACE1004130	63.87	69.03	63.79	1.08	1
PLACE1004149	177.55	168.78	207.8	0.95	1.17
PLACE1004156	114.58	150.12	132.43	1.31	1.16
PLACE1004160	244.43	317.97	314.18	1.3	1.29
PLACE1004161	21.36	26.5	25.84	1	1

(696)

特開 2002-191363

1389

1390

PLACE1004166	46.55	64.11	41.06	1.38	0.88
PLACE1004168	43.73	52.05	49.77	1.19	1.14
PLACE1004170	43.27	46.37	37.07	1.07	0.92
PLACE1004178	33.77	30.19	46.55	1	1.16
PLACE1004183	51.1	50.95	56.22	1	1.1
PLACE1004197	31.08	38.77	28.52	1	1
PLACE1004199	111.09	132.66	120.97	1.19	1.09
PLACE1004203	18.22	21.91	21.51	1	1
PLACE1004242	163.08	167.98	126.69	1.03	0.78
PLACE1004249	435.63	372.66	433.16	0.86	0.99
PLACE1004255	32.74	27.54	29.86	1	1
PLACE1004256	108	91.63	125.22	0.85	1.16
PLACE1004257	53.87	62.72	62.43	1.16	1.16
PLACE1004258	34.99	40.17	32.95	1	1
PLACE1004270	44.65	40.89	34.69	0.92	0.9
PLACE1004272	27.85	29.4	37.92	1	1
PLACE1004273	2399.9	2465.86	2118.46	1.03	0.88
PLACE1004274	92.87	93.1	85.05	1	0.92
PLACE1004277	91.27	98.55	65.14	1.08	0.71
PLACE1004279	59.07	54.01	41.5	0.91	0.7
PLACE1004282	61.13	67.16	45.79	1.1	0.75
PLACE1004284	43.87	53.56	64.09	1.22	1.46
PLACE1004289	94.78	95.17	83.09	1	0.88
PLACE1004299	58.09	235	267.04	4.05	4.6
PLACE1004302	35.49	35.83	21.22	1	1
PLACE1004305	35.37	24.85	23.42	1	1
PLACE1004316	44.19	41.05	43.26	0.93	0.98
PLACE1004322	21.55	19.73	14.91	1	1
PLACE1004325	48.04	49	55.01	1.02	1.15
PLACE1004332	69.17	53.73	54.6	0.78	0.79
PLACE1004336	120.69	133.2	92.79	1.1	0.77
PLACE1004346	24.94	22.62	24.96	1	1
PLACE1004358	60.72	45.68	58.37	0.75	0.96
PLACE1004376	199.94	252.27	215.24	1.26	1.08
PLACE1004384	60.23	55.46	45.37	0.92	0.75
PLACE1004385	16.39	12.29	13.4	1	1
PLACE1004388	31.51	33.43	37.36	1	1
PLACE1004405	26.54	39.85	27.89	1	1
PLACE1004407	166.24	172.82	137.19	1.04	0.83
PLACE1004424	26.57	21.02	21.24	1	1
PLACE1004425	60.07	75.63	53.78	1.26	0.9
PLACE1004427	65.17	71.34	56.67	1.09	0.87
PLACE1004428	57.76	61.38	49.93	1.06	0.86
PLACE1004433	117.69	111.26	87.78	0.95	0.75
PLACE1004435	46.62	46.49	64.22	1	1.38
PLACE1004437	90.85	88.51	97.35	0.97	1.07
PLACE1004441	202.84	222.16	156.5	1.1	0.77
PLACE1004446	21.83	23.05	23.83	1	1
PLACE1004450	28.93	29.69	29.16	1	1
PLACE1004451	43.89	47.63	37.97	1.09	0.91

(697)

特開 2002-191363

1391

1392

PLACE1004456	93.62	99.48	112.04	1.06	1.2
PLACE1004458	172.89	222.87	311.93	1.29	1.8
PLACE1004460	17.84	22.83	21.38	1	1
PLACE1004467	105.43	114.12	107.7	1.08	1.02
PLACE1004471	151.27	148.35	121.95	0.98	0.81
PLACE1004473	32.88	24.89	24.19	1	1
PLACE1004475	311.31	286.34	295.05	0.92	0.95
PLACE1004482	58.85	50.7	45.07	0.86	0.77
PLACE1004491	50.15	31.42	45.64	0.8	0.91
PLACE1004492	574.64	611.87	545	1.06	0.95
PLACE1004506	55.23	43.9	63.47	0.79	1.15
PLACE1004507	36.91	51.87	51.67	1.3	1.29
PLACE1004510	46.45	58.12	43.51	1.25	0.94
PLACE1004516	35.91	31.26	35.07	1	1
PLACE1004518	40.44	46.56	32.76	1.15	0.99
PLACE1004519	29.69	34.96	23.94	1	1
PLACE1004520	39.35	54.81	32.81	1.37	1
PLACE1004530	81.51	148.47	119.38	1.82	1.46
PLACE1004545	26.79	37.64	25.2	1	1
PLACE1004547	76.6	72.83	116.41	0.95	1.52
PLACE1004548	71.71	95.33	72.11	1.33	1.01
PLACE1004550	45.22	45.8	43.85	1.01	0.97
PLACE1004551	32.29	37.21	26.81	1	1
PLACE1004559	44.74	33.16	37.64	0.89	0.89
PLACE1004562	28.12	38.37	24.76	1	1
PLACE1004564	37.79	44.05	36.38	1.1	1
PLACE1004604	69.46	43.79	70.41	0.63	1.01
PLACE1004611	81.59	120.14	93.91	1.47	1.15
PLACE1004629	53.91	94.48	58.88	1.75	1.09
PLACE1004630	24.94	28.66	24.06	1	1
PLACE1004637	82.97	92.18	76.13	1.11	0.92
PLACE1004645	705.25	678.73	617.05	0.96	0.87
PLACE1004646	36.08	34.75	34.74	1	1
PLACE1004648	29.15	23.9	22.52	1	1
PLACE1004655	782.3	922.68	842.27	1.18	1.08
PLACE1004658	68.23	66.04	70.48	0.97	1.03
PLACE1004664	29.86	25.07	23.6	1	1
PLACE1004672	117.85	116.26	135.22	0.99	1.15
PLACE1004674	135.11	150.95	123.26	1.12	0.91
PLACE1004681	76.94	79.8	70.45	1.04	0.92
PLACE1004686	164.64	180.61	159.85	1.1	0.97
PLACE1004690	138.95	137.9	160.53	0.99	1.16
PLACE1004691	47.58	44.73	44.23	0.94	0.93
PLACE1004693	44.67	55.19	52.33	1.24	1.17
PLACE1004701	51.89	124.32	146.8	2.4	2.83
PLACE1004705	24.89	32.52	25.9	1	1
PLACE1004708	207.22	227.65	205.56	1.1	0.99
PLACE1004716	158.09	131.22	159.6	0.83	1.01
PLACE1004722	39.69	47.66	34.38	1.19	1
PLACE1004736	103.31	78.43	109.86	0.76	1.06

(698)

特開 2002-191363

1393

1394

PLACE1004737	37.33	40.44	27.74	1.01	1
PLACE1004740	70.48	106.24	121.63	1.51	1.73
PLACE1004743	35.03	27.05	25.32	1	1
PLACE1004751	54.43	69.55	64.09	1.28	1.18
PLACE1004757	177.06	181.12	169.77	1.02	0.96
PLACE1004761	34	31.06	35.86	1	1
PLACE1004773	30.98	28.52	30.42	1	1
PLACE1004775	14.4	16.78	10.02	1	1
PLACE1004777	26.78	34.18	24.46	1	1
PLACE1004793	30.43	36.04	28.51	1	1
PLACE1004796	46.08	51.98	46.13	1.13	1
PLACE1004804	21.4	25.53	29.44	1	1
PLACE1004813	108.55	112.81	78.88	1.04	0.73
PLACE1004814	217.28	198.39	234.28	0.91	1.08
PLACE1004815	41.79	44.51	36.77	1.07	0.96
PLACE1004816	25.64	24.95	18.01	1	1
PLACE1004824	94.12	86.16	80.67	0.92	0.86
PLACE1004827	42.22	37.37	43.88	0.95	1.04
PLACE1004836	53.01	50.84	41.19	0.96	0.78
PLACE1004838	25.58	24.86	26.19	1	1
PLACE1004840	32.92	31.37	23.93	1	1
PLACE1004842	37.6	33.39	23.96	1	1
PLACE1004850	27.87	30.65	23.65	1	1
PLACE1004868	31.05	27.28	32.01	1	1
PLACE1004885	41.35	54.34	36.85	1.31	0.97
PLACE1004886	26.38	36.64	32.21	1	1
PLACE1004887	265.33	679.13	511.75	2.56	1.93
PLACE1004896	35.15	55.19	48.06	1.38	1.2
PLACE1004900	110.48	137.06	104.42	1.24	0.95
PLACE1004902	56.18	63.04	62.72	1.12	1.12
PLACE1004904	35.56	41.39	44.64	1.03	1.12
PLACE1004911	101.01	83.57	101.41	0.83	1
PLACE1004913	33.78	28.38	26.83	1	1
PLACE1004918	28.12	30.06	20.87	1	1
PLACE1004930	44.21	39.5	43.9	0.9	0.99
PLACE1004934	30.02	33.97	36.31	1	1
PLACE1004937	41.99	43.62	36.19	1.04	0.95
PLACE1004949	387.69	536.79	490.03	1.38	1.26
PLACE1004969	30.07	28.59	21.77	1	1
PLACE1004970	17.22	12.03	17.88	1	1
PLACE1004972	37.41	41.08	40.74	1.03	1.02
PLACE1004974	22.2	26.49	24.87	1	1
PLACE1004975	25.19	26.69	21.49	1	1
PLACE1004979	99.07	105.39	101.37	1.06	1.02
PLACE1004982	87.22	92.31	86.84	1.06	1
PLACE1004985	29.21	28.18	25.76	1	1
PLACE1005003	75.58	71.91	54.65	0.95	0.72
PLACE1005004	23.89	17.83	12.04	1	1
PLACE1005005	82.75	84.73	71.73	1.02	0.87
PLACE1005011	778.5	847.92	631.14	1.09	0.81

(699)

特開 2002-191363

1395

1396

PLACE1005026	28.8	40.65	36.67	1.02	1
PLACE1005027	49.51	56.36	53.55	1.14	1.08
PLACE1005031	62.85	74.25	45.61	1.18	0.73
PLACE1005036	143.79	167.62	129.45	1.17	0.9
PLACE1005041	49.86	45.09	51.96	0.9	1.04
PLACE1005046	84.21	112.84	82.55	1.34	0.98
PLACE1005047	22.51	27.41	16.71	1	1
PLACE1005052	90.85	113.45	114.68	1.25	1.26
PLACE1005055	50.92	76.6	52.3	1.5	1.03
PLACE1005066	75.14	83.4	86.5	1.11	1.15
PLACE1005077	30.3	30.8	20.94	1	1
PLACE1005085	83.67	127.5	74.04	1.52	0.88
PLACE1005086	101.48	119.52	84.13	1.18	0.83
PLACE1005088	357.2	403.63	377.94	1.13	1.06
PLACE1005089	37.14	48.9	29.41	1.22	1
PLACE1005101	42.93	53.29	78.3	1.24	1.82
PLACE1005102	56.52	82.2	74.5	1.45	1.32
PLACE1005108	72.54	88.83	80.32	1.22	1.11
PLACE1005110	13.81	14.71	11.43	1	1
PLACE1005111	20.64	21.67	18.57	1	1
PLACE1005123	123.7	108.42	108.63	0.88	0.88
PLACE1005124	43.31	45.57	29.89	1.05	0.92
PLACE1005128	132.8	180.99	174.06	1.36	1.31
PLACE1005130	38.65	55.88	39.43	1.4	1
PLACE1005141	130.79	134.07	116.15	1.03	0.89
PLACE1005146	25.44	35.02	27.31	1	1
PLACE1005152	54.53	46.52	69.96	0.85	1.28
PLACE1005157	30.2	33.7	30.13	1	1
PLACE1005162	62.82	53.03	59.26	0.84	0.94
PLACE1005170	49.13	37.62	40.27	0.81	0.82
PLACE1005176	17.88	22.59	17.36	1	1
PLACE1005181	15.51	13.04	18.65	1	1
PLACE1005184	89.71	87.43	97.88	0.97	1.09
PLACE1005186	68.29	59.84	58.49	0.88	0.86
PLACE1005187	43.18	43.45	28.94	1.01	0.93
PLACE1005189	93.46	75.94	97.23	0.81	1.04
PLACE1005193	27.01	24.91	29.9	1	1
PLACE1005200	90.96	100.02	104.26	1.1	1.15
PLACE1005206	35.25	30.09	34.22	1	1
PLACE1005216	72.34	62.49	46.77	0.86	0.65
PLACE1005223	56.97	70.55	66.1	1.24	1.16
PLACE1005225	30.95	34.01	22.19	1	1
PLACE1005232	93.45	98.34	74.05	1.05	0.79
PLACE1005239	27.92	29.83	20.54	1	1
PLACE1005243	46.36	51.14	47.55	1.1	1.03
PLACE1005250	63.7	64.27	69.23	1.01	1.09
PLACE1005261	39.94	47.49	45.93	1.19	1.15
PLACE1005266	50.13	47.62	41.62	0.95	0.83
PLACE1005271	105.46	119.85	107.51	1.14	1.02
PLACE1005277	23.12	23.98	23.79	1	1

(700)

特開 2002-191363

1397

1398

PLACE1005287	34.49	30.55	24.59	1	1
PLACE1005299	227.75	244.53	240.7	1.07	1.06
PLACE1005305	69.35	58.03	70.36	0.84	1.01
PLACE1005307	47.35	45	36.3	0.95	0.84
PLACE1005308	69.41	60.66	61.59	0.87	0.89
PLACE1005313	30.06	30.16	21.98	1	1
PLACE1005320	23.98	27.05	34.24	1	1
PLACE1005327	115.46	121.53	98.08	1.05	0.85
PLACE1005331	30.68	23.63	27.98	1	1
PLACE1005335	21.07	19.09	21.14	1	1
PLACE1005336	59.36	64.07	46.08	1.08	0.78
PLACE1005351	214.19	250.8	232.92	1.17	1.09
PLACE1005366	40.76	25.34	38.17	0.98	0.98
PLACE1005373	39.99	49.78	40.8	1.24	1.02
PLACE1005374	72.3	70.1	60.1	0.97	0.83
PLACE1005383	28.31	37.28	28.35	1	1
PLACE1005388	20.09	15.33	18.44	1	1
PLACE1005409	41.93	44.79	33.57	1.07	0.95
PLACE1005410	88.35	94.3	106.85	1.07	1.21
PLACE1005426	19.09	18.98	20.59	1	1
PLACE1005431	88.69	108.82	135.42	1.23	1.53
PLACE1005453	71.2	88.48	54.16	1.24	0.76
PLACE1005467	77.47	67.27	64.65	0.87	0.83
PLACE1005471	25.1	22.55	17.51	1	1
PLACE1005476	18.71	20.07	18.3	1	1
PLACE1005477	41.23	32.25	32.73	0.97	0.97
PLACE1005480	17.77	14.42	12.46	1	1
PLACE1005481	50.84	54.24	40.04	1.07	0.79
PLACE1005494	20.43	21.23	13.29	1	1
PLACE1005495	40.63	117.46	154.7	2.89	3.81
PLACE1005497	42.63	75.44	65.24	1.77	1.53
PLACE1005499	80.45	84.52	88.63	1.05	1.1
PLACE1005502	27.31	28.42	23.04	1	1
PLACE1005513	23.38	27.63	25.62	1	1
PLACE1005515	17.68	23.24	12.33	1	1
PLACE1005519	52.35	24.62	32.27	0.76	0.76
PLACE1005526	20.49	19.76	12.62	1	1
PLACE1005528	99.57	107.34	90.46	1.08	0.91
PLACE1005530	107.92	99.06	99.36	0.92	0.92
PLACE1005536	27.52	29.04	23.69	1	1
PLACE1005539	22.2	17.91	26.37	1	1
PLACE1005543	39.44	35.71	33.19	1	1
PLACE1005544	43.12	40.95	42.44	0.95	0.98
PLACE1005550	66.14	57.67	48.6	0.87	0.73
PLACE1005554	23.4	23.49	18.53	1	1
PLACE1005557	36.97	33.18	33	1	1
PLACE1005563	30.06	22.35	18.23	1	1
PLACE1005569	18.5	16.08	13.5	1	1
PLACE1005574	40.77	30.21	35.7	0.98	0.98
PLACE1005584	53.92	36.99	33.41	0.74	0.74

(701)

特開 2002-191363

1399

1400

PLACE1005590	68.27	70.39	50.66	1.03	0.74
PLACE1005595	47.62	51.56	48.11	1.08	1.01
PLACE1005601	32.77	30.9	27.67	1	1
PLACE1005603	32.65	27.59	23.46	1	1
PLACE1005604	67.16	64.62	58.72	0.96	0.87
PLACE1005611	31.91	31.2	21.85	1	1
PLACE1005622	27.53	25.29	20.54	1	1
PLACE1005623	69.61	74.43	44.64	1.07	0.64
PLACE1005630	57.78	77.87	63.31	1.35	1.1
PLACE1005639	23.29	30.52	21.47	1	1
PLACE1005646	54.09	58.74	66.48	1.09	1.23
PLACE1005647	31.24	27.78	22.16	1	1
PLACE1005648	174.8	182.82	151.55	1.05	0.87
PLACE1005653	40.53	53.46	41.03	1.32	1.01
PLACE1005656	34.66	27.91	19.94	1	1
PLACE1005659	27.58	27.75	20.25	1	1
PLACE1005660	65.26	79.83	68.7	1.22	1.05
PLACE1005664	68.48	102.48	82.87	1.5	1.21
PLACE1005666	67.44	89.33	65.27	1.32	0.97
PLACE1005669	68.49	89.01	66.6	1.3	0.97
PLACE1005682	42.95	56.26	53.91	1.31	1.26
PLACE1005698	26.95	22.03	29.51	1	1
PLACE1005708	524.06	409.37	510.49	0.78	0.97
PLACE1005725	84.61	74.38	60.65	0.88	0.72
PLACE1005727	39.57	43.65	52.09	1.09	1.3
PLACE1005730	21.15	15.02	28.98	1	1
PLACE1005736	91.38	85.92	104.25	0.94	1.14
PLACE1005739	19.52	17.35	20.44	1	1
PLACE1005745	87.46	101.97	90.38	1.17	1.03
PLACE1005752	40	32.42	36.99	0.81	0.92
PLACE1005755	22.36	23.11	27.42	1	1
PLACE1005756	819	846.51	766.08	1.03	0.94
PLACE1005760	1062.4	953.29	852.14	0.9	0.8
PLACE1005763	95.45	96.07	92.3	1.01	0.97
PLACE1005768	58.28	58.42	49.18	1	0.84
PLACE1005771	65.24	83.61	81.92	1.28	1.26
PLACE1005783	56.7	49.41	41.56	0.87	0.73
PLACE1005799	84.34	62.07	66.02	0.74	0.78
PLACE1005802	40.45	36.71	22.56	0.99	0.99
PLACE1005803	52.4	52.65	55.01	1	1.05
PLACE1005804	30.97	22.7	26.77	1	1
PLACE1005813	1808.1	1743.44	1328.93	0.96	0.73
PLACE1005815	95.63	82.59	64.07	0.86	0.67
PLACE1005828	66.49	73.06	63.39	1.1	0.95
PLACE1005833	643.83	631.1	507.77	0.98	0.79
PLACE1005834	48.61	47.74	45.99	0.98	0.95
PLACE1005835	360.43	407.67	352.53	1.13	0.98
PLACE1005836	15.6	19.62	14.08	1	1
PLACE1005845	29.39	26.53	21.53	1	1
PLACE1005850	43.16	48.82	40.66	1.13	0.94

(702)

特開 2002-191363

1401

1402

PLACE1005851	24.59	29.45	23.01	1	1
PLACE1005856	31.36	36.06	38.2	1	1
PLACE1005875	28.19	27.56	23.48	1	1
PLACE1005876	38.64	32.99	39.6	1	1
PLACE1005878	28.58	30.51	22.54	1	1
PLACE1005880	32.67	30.18	23.51	1	1
PLACE1005884	26.98	25.83	20.75	1	1
PLACE1005890	35.41	31.1	33.76	1	1
PLACE1005898	102.07	112.8	93.03	1.11	0.91
PLACE1005913	94.01	97.52	77.21	1.04	0.82
PLACE1005921	32.53	33.54	19.76	1	1
PLACE1005923	19.81	19.65	16.69	1	1
PLACE1005925	35.08	37.64	17.11	1	1
PLACE1005927	20.75	20.69	18.83	1	1
PLACE1005932	24.82	24	17.52	1	1
PLACE1005934	42.12	41.03	39.84	0.97	0.95
PLACE1005936	27.74	24.68	25.54	1	1
PLACE1005939	241.35	526.2	524.44	2.18	2.17
PLACE1005951	29.76	26.23	25.81	1	1
PLACE1005953	22.65	23.03	23.62	1	1
PLACE1005955	32.99	32.9	28.43	1	1
PLACE1005966	34.05	28.76	23.82	1	1
PLACE1005968	32.82	26.94	42.21	1	1.06
PLACE1005975	56.15	93.88	70.1	1.67	1.25
PLACE1005990	17.45	27.59	15.96	1	1
PLACE1005997	978.15	2219.43	3051.68	2.27	3.12
PLACE1006002	130.64	147.49	122.1	1.13	0.93
PLACE1006003	47.98	54.08	51.82	1.13	1.08
PLACE1006011	43.98	54.19	44.58	1.23	1.01
PLACE1006017	38.94	43.81	30.38	1.1	1
PLACE1006037	26.17	24.55	38.53	1	1
PLACE1006040	37.97	39.89	41.47	1	1.04
PLACE1006063	58.1	43.53	43.45	0.75	0.75
PLACE1006071	41.95	35.58	31.87	0.95	0.95
PLACE1006073	59.05	60.62	48.62	1.03	0.82
PLACE1006074	46.55	44.08	45.19	0.95	0.97
PLACE1006076	60.92	62.26	43.74	1.02	0.72
PLACE1006079	26.25	30.84	17.99	1	1
PLACE1006093	21.09	19.72	15.98	1	1
PLACE1006116	33.38	37.02	38.15	1	1
PLACE1006119	72.16	83.88	61.88	1.16	0.86
PLACE1006129	37	24.91	26.66	1	1
PLACE1006139	103.34	92.13	64.98	0.89	0.63
PLACE1006143	77.31	82.9	55.21	1.07	0.71
PLACE1006157	73.63	59.36	39.19	0.81	0.54
PLACE1006159	50.76	58.41	56	1.15	1.1
PLACE1006164	38.42	29.2	24.79	1	1
PLACE1006167	30.05	34.28	28.7	1	1
PLACE1006170	54.12	53.86	45.69	1	0.84
PLACE1006181	51.85	61.22	75.19	1.18	1.45

(703)

特開 2002-191363

1403

1404

PLACE1006187	20.35	22.27	11.76	1	1
PLACE1006195	36.98	41.33	31.38	1.03	1
PLACE1006196	46.25	49.99	34.23	1.08	0.86
PLACE1006197	35.41	80.45	36.64	2.01	1
PLACE1006198	20.33	24.15	10.65	1	1
PLACE1006205	27.04	30.33	47.81	1	1.2
PLACE1006208	166.18	160.11	155.9	0.96	0.94
PLACE1006211	81.85	110.21	141.57	1.35	1.73
PLACE1006219	43.13	49.69	43.38	1.15	1.01
PLACE1006223	18.71	20.08	11.24	1	1
PLACE1006225	26.16	18.15	12.96	1	1
PLACE1006236	38.99	51.3	34.8	1.28	1
PLACE1006239	21.14	21.54	22.6	1	1
PLACE1006245	32.35	38.28	31.77	1	1
PLACE1006246	41.33	40.97	38.4	0.99	0.97
PLACE1006248	29.34	37.42	27.46	1	1
PLACE1006262	27.21	37.14	22.75	1	1
PLACE1006269	14.07	11.07	12.08	1	1
PLACE1006275	32.47	29.42	27.81	1	1
PLACE1006277	14.32	12.83	14.05	1	1
PLACE1006288	121.68	126.55	150.05	1.04	1.23
PLACE1006290	44.54	37.22	52.02	0.9	1.17
PLACE1006298	24.6	18.55	24.34	1	1
PLACE1006311	224.76	362.9	478.96	1.61	2.13
PLACE1006318	39.28	35.78	29.29	1	1
PLACE1006325	104.31	87.97	80.82	0.84	0.77
PLACE1006331	63.21	61.33	59.62	0.97	0.94
PLACE1006335	50.13	45.24	37.81	0.9	0.8
PLACE1006357	27.55	22.91	22.51	1	1
PLACE1006360	31.91	28.01	32.95	1	1
PLACE1006364	34.98	32.36	36.01	1	1
PLACE1006365	16.88	20.54	23.57	1	1
PLACE1006368	35.92	33.82	27.34	1	1
PLACE1006371	33.44	37.43	26.91	1	1
PLACE1006373	87.74	94.97	83.52	1.08	0.95
PLACE1006382	33.59	26.24	29.91	1	1
PLACE1006385	26.01	27.2	27.21	1	1
PLACE1006391	67.93	81.48	92.19	1.2	1.36
PLACE1006412	89.26	93.45	99.21	1.05	1.11
PLACE1006414	18.25	22.77	19.71	1	1
PLACE1006419	42.52	42.88	32.14	1.01	0.94
PLACE1006438	60	55.71	43.01	0.93	0.72
PLACE1006443	64.71	77.33	79.64	1.2	1.23
PLACE1006445	34.57	26.21	29.93	1	1
PLACE1006447	46.13	47.27	45.94	1.02	1
PLACE1006466	30.92	25.74	21.02	1	1
PLACE1006469	28.34	25.31	26.97	1	1
PLACE1006470	71.3	79.67	74.87	1.12	1.05
PLACE1006472	247.42	236.96	256.25	0.96	1.04
PLACE1006476	66.33	66.38	56.39	1	0.85

(704)

特開 2002-191363

1405

1406

PLACE1006482	55.71	67.95	39.62	1.22	0.72
PLACE1006488	263.14	243.49	211.79	0.93	0.8
PLACE1006492	74.4	85.96	60.14	1.16	0.81
PLACE1006506	39.87	39.23	42.33	1	1.06
PLACE1006515	31.54	34.21	43.45	1	1.09
PLACE1006516	28.57	28.1	32.19	1	1
PLACE1006520	50.97	51.84	36.08	1.02	0.78
PLACE1006521	91.91	93.22	93.9	1.01	1.02
PLACE1006529	77.96	70.96	85.63	0.91	1.1
PLACE1006531	17.55	16.72	17.3	1	1
PLACE1006534	28.78	26.63	23.78	1	1
PLACE1006540	97.91	113.43	90.41	1.16	0.92
PLACE1006549	24.66	26.8	21.47	1	1
PLACE1006550	30.72	30.14	34.44	1	1
PLACE1006552	25.88	21.37	17.37	1	1
PLACE1006557	20.03	20.85	22.74	1	1
PLACE1006563	32.75	38.63	32.31	1	1
PLACE1006579	38.36	50.2	53.09	1.26	1.33
PLACE1006594	1470.4	2343.7	2977.72	1.59	2.03
PLACE1006598	25.17	28.2	20.86	1	1
PLACE1006607	102.62	96.98	63.83	0.95	0.62
PLACE1006610	161.44	157.4	158.37	0.97	0.98
PLACE1006615	170.53	230.37	181.33	1.35	1.06
PLACE1006617	41.36	52.47	40.33	1.27	0.98
PLACE1006618	61.9	63.49	60.92	1.03	0.98
PLACE1006626	18.78	18.58	15.81	1	1
PLACE1006629	39.09	28.14	26.39	1	1
PLACE1006637	47.74	54.16	45.68	1.13	0.96
PLACE1006640	19.56	21.49	18.75	1	1
PLACE1006644	34.47	24.66	38.52	1	1
PLACE1006657	36.76	43.91	33.57	1.1	1
PLACE1006673	105.75	118.39	90.89	1.12	0.86
PLACE1006678	25.26	26.33	21.23	1	1
PLACE1006682	32.57	35.48	39.14	1	1
PLACE1006684	24.76	15.56	12.72	1	1
PLACE1006698	21.9	21.87	19.07	1	1
PLACE1006704	29.17	27.59	21.49	1	1
PLACE1006708	52.21	56.79	44.42	1.09	0.85
PLACE1006711	90.49	83.29	83.69	0.92	0.92
PLACE1006714	61.05	49.27	37.68	0.81	0.66
PLACE1006716	23.05	15.83	18.42	1	1
PLACE1006731	38.28	45.12	38.86	1.13	1
PLACE1006754	18.26	15.48	13.94	1	1
PLACE1006760	54.5	61.53	57.07	1.13	1.05
PLACE1006779	32.63	28.19	24.73	1	1
PLACE1006782	22.28	17.96	16.97	1	1
PLACE1006783	26.21	29.78	19.3	1	1
PLACE1006786	30.43	23.44	19.4	1	1
PLACE1006792	96.99	109.36	85.04	1.13	0.88
PLACE1006795	21.7	18.73	12.96	1	1

(705)

特開 2002-191363

1407

1408

PLACE1006800	32.54	34.66	29.4	1	1
PLACE1006805	65	91.51	86.82	1.41	1.34
PLACE1006809	55.84	58.93	56.4	1.06	1.01
PLACE1006815	38.62	40.32	33.93	1.01	1
PLACE1006819	11.29	13.41	5.54	1	1
PLACE1006820	132.74	162.76	119.25	1.23	0.9
PLACE1006826	49.88	87.77	43.48	1.76	0.87
PLACE1006829	123.27	130.22	102.14	1.06	0.83
PLACE1006853	29.17	41.16	32.7	1.03	1
PLACE1006860	17.5	23.58	21.3	1	1
PLACE1006867	66.64	94.76	67.34	1.42	1.01
PLACE1006875	43.58	44.55	38.5	1.02	0.92
PLACE1006878	49.53	49.03	45.07	0.99	0.91
PLACE1006883	76.12	68.45	71.78	0.9	0.94
PLACE1006898	46.07	40.11	68.46	0.87	1.49
PLACE1006901	31.34	33.46	37.51	1	1
PLACE1006904	58.07	68.41	69.27	1.18	1.19
PLACE1006917	60.31	46.57	73.27	0.77	1.21
PLACE1006932	25.98	21.08	20.62	1	1
PLACE1006935	14.99	15.43	17.56	1	1
PLACE1006956	33.28	22.73	21.83	1	1
PLACE1006958	26.5	20.75	19.07	1	1
PLACE1006959	67.11	66.82	64.38	1	0.96
PLACE1006961	90.14	97.32	90.46	1.08	1
PLACE1006962	92.89	89.3	88.91	0.96	0.96
PLACE1006966	35.53	38.49	39.19	1	1
PLACE1006979	25.57	36.6	24.14	1	1
PLACE1006989	29.39	43.87	33.32	1.1	1
PLACE1007001	64.69	60.54	57.94	0.94	0.9
PLACE1007014	34.47	27.4	25.01	1	1
PLACE1007021	37.43	35.79	26.42	1	1
PLACE1007026	33.55	27.29	38.92	1	1
PLACE1007028	70.75	79.26	68.52	1.12	0.97
PLACE1007038	905.48	1441.98	1352.28	1.59	1.49
PLACE1007040	42	29.8	29.82	0.95	0.95
PLACE1007045	36.67	50.06	37.33	1.25	1
PLACE1007048	679.34	713.4	741.8	1.05	1.09
PLACE1007053	31.94	30.93	29.32	1	1
PLACE1007068	860.62	1042.12	659.27	1.21	0.77
PLACE1007070	71.9	54.02	58.47	0.75	0.81
PLACE1007076	146.49	190.93	165.54	1.3	1.13
PLACE1007077	32.04	29.37	39.47	1	1
PLACE1007081	29.72	27.35	21.94	1	1
PLACE1007082	23.97	23.42	30.77	1	1
PLACE1007092	69.29	69.41	61.52	1	0.89
PLACE1007096	21.59	22.38	18.71	1	1
PLACE1007097	29.53	26.42	24.05	1	1
PLACE1007099	31.63	31.56	22.21	1	1
PLACE1007105	20	18.95	19.48	1	1
PLACE1007108	38.02	33.29	48.92	1	1.22

(706)

特開 2002-191363

1409

1410

PLACE1007111	23.6	19.67	22.41	1	1
PLACE1007112	19.93	23.21	22.24	1	1
PLACE1007130	24.68	18.62	16.94	1	1
PLACE1007132	57.55	73.92	47.55	1.28	0.83
PLACE1007140	18.39	18.58	14.36	1	1
PLACE1007143	35.84	33.62	34.44	1	1
PLACE1007169	34.92	47.03	50.92	1.18	1.27
PLACE1007178	19.73	19.53	17.73	1	1
PLACE1007190	28	37.16	35.42	1	1
PLACE1007201	18.63	16.17	17.55	1	1
PLACE1007202	677.92	712.91	790.84	1.05	1.17
PLACE1007226	24.8	33.84	26.76	1	1
PLACE1007238	33.43	38.99	29.72	1	1
PLACE1007239	20.1	18.41	23.99	1	1
PLACE1007242	16.1	16.07	12.84	1	1
PLACE1007243	24.97	28.75	26.19	1	1
PLACE1007247	51.59	68.41	90.72	1.33	1.76
PLACE1007257	42.7	49.42	38.19	1.16	0.94
PLACE1007274	69.44	86.8	65.54	1.25	0.94
PLACE1007276	48.55	59.63	39.58	1.23	0.82
PLACE1007282	43.99	53.32	49.34	1.21	1.12
PLACE1007286	134.31	145.39	129.15	1.08	0.96
PLACE1007296	123.21	122.62	140.64	1	1.14
PLACE1007301	28.54	25.3	16.19	1	1
PLACE1007314	44.52	56.15	56.98	1.26	1.28
PLACE1007317	27.02	28.27	23.95	1	1
PLACE1007329	37.74	46.99	36.74	1.17	1
PLACE1007338	74.91	71.97	76.65	0.96	1.02
PLACE1007342	23.57	21.57	14.99	1	1
PLACE1007345	23.84	16.63	15.18	1	1
PLACE1007346	63.49	71.76	66.56	1.13	1.05
PLACE1007359	28.39	32.41	26.65	1	1
PLACE1007367	159.7	186.86	156.38	1.17	0.98
PLACE1007375	19.55	21.28	14.79	1	1
PLACE1007377	32.15	26.06	26.24	1	1
PLACE1007386	51.65	39.6	35.88	0.77	0.77
PLACE1007392	57.87	39.48	53.45	0.69	0.92
PLACE1007402	24.24	21.89	19.65	1	1
PLACE1007409	26.87	31.02	20.7	1	1
PLACE1007416	42.3	50.8	46.06	1.2	1.09
PLACE1007420	157.5	187.73	170.92	1.19	1.09
PLACE1007431	53.07	48.14	51.76	0.91	0.98
PLACE1007450	57.18	56.14	36.36	0.98	0.7
PLACE1007452	47.38	52.61	44.05	1.11	0.93
PLACE1007454	808.25	1172.33	677.25	1.45	0.84
PLACE1007460	22.94	51.42	18.14	1.29	1
PLACE1007478	23.46	29.22	21.9	1	1
PLACE1007484	32.11	29.62	26.58	1	1
PLACE1007488	23.81	42.75	27.1	1.07	1
PLACE1007507	32.1	37.71	39.41	1	1

(707)

特開 2002-191363

1411

1412

PLACE1007511	15.76	5.91	8.15	1	1
PLACE1007513	161.99	171.44	146.12	1.06	0.9
PLACE1007524	34.31	42.75	22.87	1.07	1
PLACE1007525	33.3	38.5	31.23	1	1
PLACE1007537	522.32	806.81	756.73	1.54	1.45
PLACE1007544	45.92	59.57	40.41	1.3	0.88
PLACE1007547	12.49	11.54	10.46	1	1
PLACE1007557	26.79	35.05	25.91	1	1
PLACE1007560	85.37	83.75	89.1	0.98	1.04
PLACE1007565	13.72	11.48	11.24	1	1
PLACE1007580	15.6	17.77	13.42	1	1
PLACE1007583	13.03	11.45	11.12	1	1
PLACE1007591	23.16	24.73	22.04	1	1
PLACE1007598	34.82	41.77	40.76	1.04	1.02
PLACE1007610	15.71	17.8	14.41	1	1
PLACE1007618	16.96	14.76	19.7	1	1
PLACE1007621	49.66	63.38	47.91	1.28	0.96
PLACE1007626	206.41	191.62	205.95	0.93	1
PLACE1007632	25.71	33.13	22.35	1	1
PLACE1007635	28.76	44.58	32.88	1.11	1
PLACE1007645	66.33	65.83	72.06	0.99	1.09
PLACE1007649	29.1	34.68	30.47	1	1
PLACE1007659	90.42	97.9	99.72	1.08	1.1
PLACE1007669	49.54	52.01	42.57	1.05	0.86
PLACE1007677	63.65	59.05	51.58	0.93	0.81
PLACE1007688	17.33	19.31	18.55	1	1
PLACE1007690	36.16	45.58	38.69	1.14	1
PLACE1007697	19.44	27.63	18.31	1	1
PLACE1007702	45.14	50.45	48.08	1.12	1.07
PLACE1007705	24.22	28.63	23.9	1	1
PLACE1007706	30.37	61.24	54.39	1.53	1.36
PLACE1007725	24.61	34.48	29.15	1	1
PLACE1007729	34.14	33.58	29.39	1	1
PLACE1007730	39.03	48.12	36.63	1.2	1
PLACE1007737	66.18	62.59	60.25	0.95	0.91
PLACE1007743	29.65	29.57	22.91	1	1
PLACE1007746	39.86	55.92	40.38	1.4	1.01
PLACE1007753	45.77	33.19	32.84	0.87	0.87
PLACE1007769	19.02	23.08	24.56	1	1
PLACE1007780	27.07	28.18	15.2	1	1
PLACE1007791	47.86	33.55	52.91	0.84	1.11
PLACE1007807	33.91	38.21	30.23	1	1
PLACE1007810	39.2	33.89	26.94	1	1
PLACE1007814	46.48	47.22	43.65	1.02	0.94
PLACE1007828	22.07	33.44	22.21	1	1
PLACE1007829	48.99	63.25	55.56	1.29	1.13
PLACE1007841	53.22	51.05	50.01	0.96	0.94
PLACE1007842	26.23	28.98	21.18	1	1
PLACE1007843	25.1	23.19	23.9	1	1
PLACE1007845	43.72	36.46	48.07	0.91	1.1

(708)

特開 2002-191363

1413

1414

PLACE1007846	28.86	32.03	27.2	1	1
PLACE1007848	29.19	38.01	26.21	1	1
PLACE1007852	32.2	42.4	27.49	1.06	1
PLACE1007858	48.92	70.66	50.24	1.44	1.03
PLACE1007866	332.62	421.67	403.75	1.27	1.21
PLACE1007871	133.08	126.7	134.77	0.95	1.01
PLACE1007877	41.18	51.83	34.76	1.26	0.97
PLACE1007878	126.48	142.52	119.91	1.13	0.95
PLACE1007881	19.21	19.92	18.79	1	1
PLACE1007885	45.12	41.68	42.69	0.92	0.95
PLACE1007897	24.63	23.73	16.27	1	1
PLACE1007908	63.89	82.77	86.71	1.3	1.36
PLACE1007922	66.14	60.47	54.63	0.91	0.83
PLACE1007946	24.76	34.34	26.2	1	1
PLACE1007950	89.89	101.23	70.71	1.13	0.79
PLACE1007954	22.84	24.75	19.71	1	1
PLACE1007955	37.84	43.08	42.66	1.08	1.07
PLACE1007956	23.21	22.74	25.64	1	1
PLACE1007958	12.85	11.41	9.59	1	1
PLACE1007965	42.59	44.77	32.57	1.05	0.94
PLACE1007969	77.66	66.93	64.71	0.86	0.83
PLACE1007971	32.43	45.19	31.39	1.13	1
PLACE1007990	38.68	34.43	34.99	1	1
PLACE1008000	20.44	19.44	11.18	1	1
PLACE1008002	22.17	33.99	22.75	1	1
PLACE1008037	16.65	21.52	14.26	1	1
PLACE1008044	37.14	25.78	26.62	1	1
PLACE1008045	36.38	23.24	30.44	1	1
PLACE1008080	29.7	29.28	28.28	1	1
PLACE1008092	24.71	24.17	18.05	1	1
PLACE1008095	34.48	41.13	26.09	1.03	1
PLACE1008105	143.94	109.41	100.26	0.76	0.7
PLACE1008107	19.18	21.02	17.6	1	1
PLACE1008111	29.75	32.08	29.35	1	1
PLACE1008113	122.11	147.06	134.03	1.2	1.1
PLACE1008122	16.38	26.8	19.23	1	1
PLACE1008129	64.74	50.69	63.95	0.78	0.99
PLACE1008132	57.78	63.8	38.67	1.1	0.69
PLACE1008137	32.83	28.72	29.18	1	1
PLACE1008174	68.35	62.42	52.45	0.91	0.77
PLACE1008177	22.86	29.48	15	1	1
PLACE1008181	21.43	22.51	16.17	1	1
PLACE1008195	34.63	40.85	43	1.02	1.08
PLACE1008198	29.8	35.31	22.76	1	1
PLACE1008201	29.51	41.32	24.42	1.03	1
PLACE1008209	40.18	38.31	21.63	1	1
PLACE1008226	100.46	120.86	72.39	1.2	0.72
PLACE1008227	65.63	79.11	45.82	1.21	0.7
PLACE1008231	20.91	25.23	20.05	1	1
PLACE1008238	43.7	29.4	37.16	0.92	0.92

(709)

特開 2002-191363

1415

1416

PLACE1008244	24.28	34.12	24.16	1	1
PLACE1008249	39.57	40.74	20.42	1.02	1
PLACE1008266	192.68	238.99	221.82	1.24	1.15
PLACE1008273	45.14	34.93	35.3	0.89	0.89
PLACE1008275	29.89	24.58	30.57	1	1
PLACE1008280	31.84	26.17	33.13	1	1
PLACE1008282	46.39	42.02	55.69	0.91	1.2
PLACE1008297	27.1	23.76	23.37	1	1
PLACE1008303	36.02	27.9	38.59	1	1
PLACE1008309	12.82	20.09	21.46	1	1
PLACE1008315	44.1	71.28	50.84	1.62	1.15
PLACE1008329	47.26	31.28	40.19	0.85	0.85
PLACE1008330	56.78	50.96	52.48	0.9	0.92
PLACE1008331	36.09	39.1	38.37	1	1
PLACE1008351	46.98	54.09	52.87	1.15	1.13
PLACE1008356	37.43	38.51	35.78	1	1
PLACE1008359	25.5	36.61	27.97	1	1
PLACE1008368	33.84	39.35	44.68	1	1.12
PLACE1008369	26.62	38.18	22.53	1	1
PLACE1008392	39.26	40.96	41.66	1.02	1.04
PLACE1008394	64.15	72.39	65.77	1.13	1.03
PLACE1008398	117.87	78.09	88.46	0.66	0.75
PLACE1008401	74.59	80.4	77.01	1.08	1.03
PLACE1008402	46.76	45.46	31	0.97	0.86
PLACE1008405	294.38	355.09	299.1	1.21	1.02
PLACE1008409	49.91	58.05	61.69	1.16	1.24
PLACE1008420	36.85	25.34	18.93	1	1
PLACE1008424	31.93	31.73	34.57	1	1
PLACE1008426	128.65	104.91	110.1	0.82	0.86
PLACE1008429	26.12	36.14	18.71	1	1
PLACE1008430	31.14	27.16	21.39	1	1
PLACE1008437	27.95	26.66	26.19	1	1
PLACE1008453	19.74	23.38	20.36	1	1
PLACE1008454	59.11	47.29	55.88	0.8	0.95
PLACE1008455	104.77	114.22	86.16	1.09	0.82
PLACE1008457	105.89	110.43	74.5	1.04	0.7
PLACE1008465	19.49	16.22	15.38	1	1
PLACE1008469	49.11	61.84	63.6	1.26	1.3
PLACE1008488	25.09	39.82	22.71	1	1
PLACE1008519	43.53	37.61	28.61	0.92	0.92
PLACE1008524	24.48	26.12	28.78	1	1
PLACE1008531	34.75	30.13	30.84	1	1
PLACE1008532	53.96	74.15	45.05	1.37	0.83
PLACE1008533	36.34	37.3	30.27	1	1
PLACE1008542	24.41	34.57	31.42	1	1
PLACE1008549	28.54	28.54	20.08	1	1
PLACE1008560	18.37	27.73	27.3	1	1
PLACE1008567	48.98	39.86	41.78	0.82	0.85
PLACE1008568	28.33	33.4	37.15	1	1
PLACE1008569	36.53	70.51	67.9	1.76	1.7

(710)

特開 2002-191363

1417

1418

PLACE1008584	33.63	55.99	53.83	1.4	1.35
PLACE1008585	58.8	75.59	59.76	1.29	1.02
PLACE1008603	581.28	561.11	342.21	0.97	0.59
PLACE1008621	24.79	24.25	20.06	1	1
PLACE1008625	11.5	11.09	11.98	1	1
PLACE1008626	16.22	13.49	12.11	1	1
PLACE1008627	29.28	30.91	37.43	1	1
PLACE1008629	65.18	56.54	148.28	0.87	2.27
PLACE1008630	22.18	21.25	23.79	1	1
PLACE1008643	34.29	35.04	29.44	1	1
PLACE1008650	14.74	15.41	13.33	1	1
PLACE1008657	55.09	57.36	43.89	1.04	0.8
PLACE1008664	23.34	23.64	24.27	1	1
PLACE1008693	43.13	41.51	30.19	0.96	0.93
PLACE1008696	31.94	34.65	25.43	1	1
PLACE1008715	47.85	46.82	42.88	0.98	0.9
PLACE1008716	41.71	53.61	31.3	1.29	0.96
PLACE1008722	54.61	45.03	44.82	0.82	0.82
PLACE1008738	88.36	84.43	83.28	0.96	0.94
PLACE1008742	162.86	160.2	161.81	0.98	0.99
PLACE1008744	14.14	22.94	12.18	1	1
PLACE1008748	22.27	24.48	14.44	1	1
PLACE1008757	34.6	42.71	27.87	1.07	1
PLACE1008766	32.11	27.78	28.26	1	1
PLACE1008785	49.07	59.73	41.43	1.22	0.84
PLACE1008790	33.27	33.02	29.64	1	1
PLACE1008798	63.72	66.89	67.61	1.05	1.06
PLACE1008807	22.01	22.03	19.94	1	1
PLACE1008808	32.24	18.82	19.03	1	1
PLACE1008813	29.43	35.83	35.39	1	1
PLACE1008836	79.18	55.11	47.12	0.7	0.6
PLACE1008851	115.23	81.63	55.52	0.71	0.48
PLACE1008854	20.39	26.55	11.71	1	1
PLACE1008864	48.15	54.26	72.45	1.13	1.5
PLACE1008867	56.85	58.24	33.04	1.02	0.7
PLACE1008876	549.37	600.39	469.11	1.09	0.85
PLACE1008887	41.83	58.22	24.8	1.39	0.96
PLACE1008902	25.21	36.34	34.18	1	1
PLACE1008911	55.54	65.78	52.58	1.18	0.95
PLACE1008917	19.14	25.45	24.12	1	1
PLACE1008920	11.29	12.12	6.23	1	1
PLACE1008925	13.83	16.73	10.48	1	1
PLACE1008930	27.53	23.49	18.05	1	1
PLACE1008934	28.64	34.88	18.8	1	1
PLACE1008941	17.54	24.91	22.06	1	1
PLACE1008947	37.22	51.14	43.96	1.28	1.1
PLACE1008984	23.24	34.33	21.6	1	1
PLACE1008985	19.99	29.03	21.28	1	1
PLACE1008994	10.53	10.54	7.43	1	1
PLACE1009020	20.44	34.43	21.28	1	1

(711)

特開 2002-191363

1419

1420

PLACE1009027	14.33	10.59	13.29	1	1
PLACE1009039	12.51	12.97	9.67	1	1
PLACE1009045	25.26	34.62	28.65	1	1
PLACE1009048	11.07	6.67	9.57	1	1
PLACE1009050	16.3	17.99	13.91	1	1
PLACE1009060	52.29	76.48	76.13	1.46	1.46
PLACE1009067	63.86	53.76	34.08	0.84	0.63
PLACE1009071	54.23	73.27	59.81	1.35	1.1
PLACE1009090	29.1	39.11	39.61	1	1
PLACE1009091	739.81	739.41	547.06	1	0.74
PLACE1009094	19.97	28.69	21.59	1	1
PLACE1009099	36.64	45.19	48.83	1.13	1.22
PLACE1009110	20.35	18.46	20.9	1	1
PLACE1009111	18.51	23.09	18.16	1	1
PLACE1009113	53.98	42.76	37.35	0.79	0.74
PLACE1009130	41.39	52.31	43.13	1.26	1.04
PLACE1009150	39.25	37.55	32.77	1	1
PLACE1009155	69.35	94.89	74.07	1.37	1.07
PLACE1009158	21.91	26.54	18.92	1	1
PLACE1009166	50.25	51.72	46.79	1.03	0.93
PLACE1009172	22.82	29.53	36.75	1	1
PLACE1009174	34.79	36.37	30.96	1	1
PLACE1009183	94.37	89.34	64.66	0.95	0.69
PLACE1009186	20.43	20.53	17.56	1	1
PLACE1009190	20.48	35.41	18.66	1	1
PLACE1009196	54.35	52.71	38.66	0.97	0.74
PLACE1009200	58.25	76.13	58.44	1.31	1
PLACE1009217	42.05	46.24	41.71	1.1	0.99
PLACE1009230	144.11	204.77	153.75	1.42	1.07
PLACE1009236	24.31	27.66	29.72	1	1
PLACE1009246	158.99	139.66	83.91	0.88	0.53
PLACE1009265	66.52	42.3	71.65	0.64	1.08
PLACE1009279	34.59	34.51	23.57	1	1
PLACE1009298	79.99	77.85	81.83	0.97	1.02
PLACE1009308	33.45	27.01	16.14	1	1
PLACE1009319	65.4	73.56	77.72	1.12	1.19
PLACE1009328	49.82	47.34	43.67	0.95	0.88
PLACE1009335	19.74	18.97	16.42	1	1
PLACE1009338	46.74	52.39	38.16	1.12	0.86
PLACE1009344	19.93	16.12	20.93	1	1
PLACE1009355	147.61	162.58	173.91	1.1	1.18
PLACE1009368	17.87	20.22	16.8	1	1
PLACE1009375	22.61	32.24	29.19	1	1
PLACE1009388	49.9	74.96	45.01	1.5	0.9
PLACE1009398	35.17	53.79	43.1	1.34	1.08
PLACE1009404	34.21	46.04	38.16	1.15	1
PLACE1009410	26.75	35.99	28.94	1	1
PLACE1009417	58.32	67.04	42.72	1.15	0.73
PLACE1009424	34.63	42.13	42.37	1.05	1.06
PLACE1009434	24.85	24.58	24.73	1	1

(712)

特開 2002-191363

1421

1422

PLACE1009443	47.53	48.97	35.82	1.03	0.84
PLACE1009444	31.89	50.08	34.61	1.25	1
PLACE1009459	18.43	17.86	18.01	1	1
PLACE1009460	67.87	55.15	59.2	0.81	0.87
PLACE1009468	61.44	73.54	56.01	1.2	0.91
PLACE1009476	18.42	23.28	12.77	1	1
PLACE1009477	38.48	38.41	40.05	1	1
PLACE1009493	14.61	13.38	11.75	1	1
PLACE1009502	24.2	18.19	17.23	1	1
PLACE1009524	18.39	17.32	15.6	1	1
PLACE1009527	34.8	37.6	27.86	1	1
PLACE1009531	202.28	205.54	217.56	1.02	1.08
PLACE1009535	41.74	48.48	38.26	1.16	0.96
PLACE1009539	52.88	57.83	52.55	1.09	0.99
PLACE1009540	74.03	82.77	96.82	1.12	1.31
PLACE1009542	20.3	17.51	18.66	1	1
PLACE1009546	26.64	16.94	13.4	1	1
PLACE1009556	19.56	22.58	16.96	1	1
PLACE1009569	42.87	51.18	43.48	1.19	1.01
PLACE1009571	28.33	26.69	25.34	1	1
PLACE1009573	55.23	54.49	56.76	0.99	1.03
PLACE1009576	43.33	35.6	25.18	0.92	0.92
PLACE1009580	66.98	54.33	52.85	0.81	0.79
PLACE1009581	35.02	40.73	37.32	1.02	1
PLACE1009587	21.2	22.71	17.17	1	1
PLACE1009593	69.23	95.74	64.2	1.38	0.93
PLACE1009595	67.55	60.93	55.12	0.9	0.82
PLACE1009596	31.43	27.75	24.05	1	1
PLACE1009600	76.24	87.08	68	1.14	0.89
PLACE1009604	51.17	39.11	25.77	0.78	0.78
PLACE1009607	105.68	136.82	84.46	1.29	0.8
PLACE1009613	27.91	34.76	27.43	1	1
PLACE1009621	70.93	83.59	74.98	1.18	1.06
PLACE1009622	51.58	101.79	73.07	1.97	1.42
PLACE1009624	36.65	51.5	30.53	1.29	1
PLACE1009637	14.3	19.67	15.18	1	1
PLACE1009639	60.81	23.05	18.6	0.66	0.66
PLACE1009654	29.81	37.96	30.06	1	1
PLACE1009659	49.17	66.83	38.29	1.36	0.81
PLACE1009665	36.25	41.13	28.7	1.03	1
PLACE1009669	20.88	23.64	26.93	1	1
PLACE1009670	27.37	30.25	30.27	1	1
PLACE1009708	24.07	37.18	35.56	1	1
PLACE1009721	28.98	18.31	19.77	1	1
PLACE1009731	35.52	28.93	25.46	1	1
PLACE1009735	40.83	32.49	55.08	0.98	1.35
PLACE1009737	35.2	30.66	37.3	1	1
PLACE1009741	18.21	26.46	34.29	1	1
PLACE1009752	40.61	30.49	23.87	0.98	0.98
PLACE1009763	49.08	51.16	54.68	1.04	1.11

(713)

特開 2002-191363

1423

1424

PLACE1009766	41.6	36.03	39.03	0.96	0.96
PLACE1009772	17.1	38.53	32.16	1	1
PLACE1009782	44.04	32.84	31.11	0.91	0.91
PLACE1009794	37.6	36.48	30.69	1	1
PLACE1009798	43.74	40.97	39.06	0.94	0.91
PLACE1009845	17.16	29.15	19.05	1	1
PLACE1009849	18.1	25.14	34.22	1	1
PLACE1009857	24.47	21.19	22.73	1	1
PLACE1009861	157.04	147.3	184.61	0.94	1.18
PLACE1009872	183.78	532.08	595.65	2.9	3.24
PLACE1009877	88.74	127.12	82	1.43	0.92
PLACE1009879	50.78	61.76	51.43	1.22	1.01
PLACE1009886	26.47	30.9	26.46	1	1
PLACE1009888	40.97	27.49	37.88	0.98	0.98
PLACE1009908	38.76	52.68	31.63	1.32	1
PLACE1009919	72.43	157.7	91.06	2.18	1.26
PLACE1009921	33.92	34.44	25.56	1	1
PLACE1009923	30.5	44.49	36.52	1.11	1
PLACE1009924	99.41	54.1	55.42	0.54	0.56
PLACE1009925	35.65	20.58	22.27	1	1
PLACE1009931	90.33	98.46	76.54	1.09	0.85
PLACE1009935	25.18	24.01	15.51	1	1
PLACE1009947	27.03	27.39	20.09	1	1
PLACE1009961	24.51	25.25	21.83	1	1
PLACE1009971	24.03	25.77	25.6	1	1
PLACE1009982	18.22	39.31	15.49	1	1
PLACE1009992	9.19	11.11	8.64	1	1
PLACE1009995	63.3	51.96	56.69	0.82	0.9
PLACE1009997	43.45	50.16	41.93	1.15	0.97
PLACE1010002	32.44	23.09	16.62	1	1
PLACE1010011	25.99	24.02	19.6	1	1
PLACE1010013	15.52	18.47	13.26	1	1
PLACE1010021	24.58	36.88	37.14	1	1
PLACE1010023	43.52	60.71	61.14	1.39	1.4
PLACE1010031	52.61	51.96	38.23	0.99	0.76
PLACE1010039	19.95	22.68	18.66	1	1
PLACE1010045	36.67	47.27	35.84	1.18	1
PLACE1010053	21.58	20.14	20.04	1	1
PLACE1010060	38.71	56.37	32.78	1.41	1
PLACE1010069	19.39	23.73	18.91	1	1
PLACE1010070	20.01	20.72	21.19	1	1
PLACE1010074	165.08	128.25	150.68	0.78	0.91
PLACE1010076	230.47	311.86	258.06	1.35	1.12
PLACE1010078	31.79	36.25	37	1	1
PLACE1010081	76.83	74.34	52.61	0.97	0.68
PLACE1010083	17.09	21.13	24.34	1	1
PLACE1010089	20.95	24.75	17.3	1	1
PLACE1010096	70.55	70.61	54.95	1	0.78
PLACE1010102	39.64	55.68	62.77	1.39	1.57
PLACE1010105	47.44	49.6	48.65	1.05	1.03

(714)

特開 2002-191363

1425

1426

PLACE1010106	40.34	52.34	43.81	1.3	1.09
PLACE1010130	40.29	44.66	30.55	1.11	0.99
PLACE1010132	56.7	85.85	65.52	1.51	1.16
PLACE1010134	20.51	15.99	20.46	1	1
PLACE1010139	415.41	498.57	421.38	1.2	1.01
PLACE1010148	39.47	45.32	27.49	1.13	1
PLACE1010152	29.32	28.44	29.8	1	1
PLACE1010155	330.96	335.95	408.91	1.02	1.24
PLACE1010156	356.21	422.83	469.57	1.19	1.32
PLACE1010161	37.57	44.88	36.68	1.12	1
PLACE1010181	34.13	35.33	27.84	1	1
PLACE1010194	24.35	22.48	25.49	1	1
PLACE1010202	18.23	20.58	13.78	1	1
PLACE1010231	31.51	46.67	24.45	1.17	1
PLACE1010235	45.66	38.3	32.66	0.88	0.88
PLACE1010237	15.65	11.4	11.73	1	1
PLACE1010251	45.81	45.52	23.23	0.99	0.87
PLACE1010261	32.21	38.4	27.38	1	1
PLACE1010270	25.56	34.7	21.54	1	1
PLACE1010273	20.85	22.99	16.74	1	1
PLACE1010274	54.13	47.79	55.26	0.88	1.02
PLACE1010277	71.1	82.25	73.8	1.16	1.04
PLACE1010293	66.72	98.76	53.8	1.48	0.81
PLACE1010297	190.1	248.33	202.48	1.31	1.07
PLACE1010300	45.22	39.59	27.21	0.88	0.88
PLACE1010310	424.98	529.37	453.64	1.25	1.07
PLACE1010321	55.25	72.06	55.13	1.3	1
PLACE1010324	25.61	25.41	17.98	1	1
PLACE1010329	50.18	65.57	47.53	1.31	0.95
PLACE1010330	55.1	76.67	55.02	1.39	1
PLACE1010335	77.42	150.11	159.9	1.94	2.07
PLACE1010341	14.28	17.1	15.38	1	1
PLACE1010342	9.17	10.68	6.34	1	1
PLACE1010346	25	30.67	21.84	1	1
PLACE1010362	35.78	48.63	28.59	1.22	1
PLACE1010364	19.36	14.8	13.85	1	1
PLACE1010368	32.4	36.5	44.4	1	1.11
PLACE1010373	55.73	66.05	70.87	1.19	1.27
PLACE1010383	105.68	48.05	71.29	0.45	0.67
PLACE1010385	6.58	5.18	2.96	1	1
PLACE1010389	149.57	123.41	125.82	0.83	0.84
PLACE1010401	15.79	15.26	16.26	1	1
PLACE1010410	28.15	37.81	34.22	1	1
PLACE1010418	36.95	33.69	39.07	1	1
PLACE1010425	16.45	18.69	17.64	1	1
PLACE1010443	273.44	343.89	242.35	1.26	0.89
PLACE1010445	32.75	43.52	36.7	1.09	1
PLACE1010481	19.87	17.5	20.62	1	1
PLACE1010482	24.01	33.97	24.55	1	1
PLACE1010491	35.66	27.01	27.99	1	1

(715)

特開 2002-191363

1427

1428

PLACE1010492	97.73	111.42	92.48	1.14	0.95
PLACE1010509	28.13	32.42	36.16	1	1
PLACE1010518	98.12	123.6	114.12	1.26	1.16
PLACE1010522	100.54	105.4	126.63	1.05	1.26
PLACE1010529	32.36	59.45	68.71	1.49	1.72
PLACE1010547	21.89	34.68	16.06	1	1
PLACE1010560	29.41	35.2	34.06	1	1
PLACE1010562	23.25	30	23.72	1	1
PLACE1010579	38.25	33.08	29.89	1	1
PLACE1010580	37.66	33.68	35.77	1	1
PLACE1010599	92.73	121.5	90.34	1.31	0.97
PLACE1010606	30.04	34.79	22.32	1	1
PLACE1010616	46.86	69.11	81.69	1.47	1.74
PLACE1010622	28.24	38.42	25.03	1	1
PLACE1010624	51.97	49.53	49.27	0.95	0.95
PLACE1010628	40.93	49.51	35.73	1.21	0.98
PLACE1010629	35.56	41.65	35.77	1.04	1
PLACE1010630	44.56	38.38	46.6	0.9	1.05
PLACE1010631	27.36	20.75	17.51	1	1
PLACE1010651	233.15	363.62	282.49	1.56	1.21
PLACE1010661	22.85	27.37	31.66	1	1
PLACE1010662	26.01	24.27	21.39	1	1
PLACE1010668	190.83	153.29	181.8	0.8	0.95
PLACE1010702	59.8	61.23	55.06	1.02	0.92
PLACE1010709	513.52	534.45	611.18	1.04	1.19
PLACE1010713	60.05	51.29	72.19	0.85	1.2
PLACE1010714	19.13	16.92	16.72	1	1
PLACE1010716	26.71	33.85	35.18	1	1
PLACE1010717	42.47	35.58	37.32	0.94	0.94
PLACE1010720	229.86	278.82	183.49	1.21	0.8
PLACE1010739	32.44	33.25	22.11	1	1
PLACE1010743	22.46	29.9	23.34	1	1
PLACE1010752	18.27	19.55	21.76	1	1
PLACE1010761	104.86	122.36	130.93	1.17	1.25
PLACE1010771	45.25	59.5	57.83	1.31	1.28
PLACE1010784	20.56	18.65	15.58	1	1
PLACE1010786	40.55	56.08	27.24	1.38	0.99
PLACE1010789	27.09	24.19	22.66	1	1
PLACE1010800	34.72	39.52	36.8	1	1
PLACE1010802	35.51	47.64	36.65	1.19	1
PLACE1010811	21.55	29.77	27	1	1
PLACE1010813	1287.7	1414.73	856.15	1.1	0.66
PLACE1010827	23	21.43	17.87	1	1
PLACE1010833	33.54	35.98	30.25	1	1
PLACE1010839	72	85.27	69.36	1.18	0.96
PLACE1010856	86.05	87.34	90.48	1.01	1.05
PLACE1010857	57.56	61.88	60.74	1.08	1.06
PLACE1010870	40.96	50.46	29.36	1.23	0.98
PLACE1010877	46.38	39.36	49.51	0.86	1.07
PLACE1010882	16.02	15.31	13.01	1	1

(716)

特開 2002-191363

1429

1430

PLACE1010891	16.64	15.08	17.9	1	1
PLACE1010896	30.16	23.93	23.2	1	1
PLACE1010900	323.44	310.25	341.11	0.96	1.05
PLACE1010916	20.94	23.68	16.14	1	1
PLACE1010917	8.78	12.15	5.58	1	1
PLACE1010924	28.2	31.02	20.59	1	1
PLACE1010925	59.4	63.62	79.68	1.07	1.34
PLACE1010926	26.05	25.68	21.29	1	1
PLACE1010942	73.09	63.9	54.96	0.87	0.75
PLACE1010943	114.11	144.64	114	1.27	1
PLACE1010944	34.71	34.89	33.54	1	1
PLACE1010947	50.97	56.41	46.75	1.11	0.92
PLACE1010954	179.34	191.9	149.28	1.07	0.83
PLACE1010960	63.89	65.16	57.38	1.02	0.9
PLACE1010965	63.35	54.39	57.39	0.86	0.91
PLACE1010968	20	25	20.64	1	1
PLACE1010978	52.43	69.4	55.7	1.32	1.06
PLACE1010982	40.37	51.27	37.65	1.27	0.99
PLACE1010990	34.79	33.81	43.07	1	1.08
PLACE1011017	114.63	113.45	90.55	0.99	0.79
PLACE1011019	66.26	57.22	82.98	0.86	1.25
PLACE1011026	61.83	68.6	53.2	1.11	0.86
PLACE1011032	44.7	54.48	36.43	1.22	0.89
PLACE1011041	31.57	35.69	21.03	1	1
PLACE1011045	35.8	33.36	50.66	1	1.27
PLACE1011046	34.53	36.4	27.94	1	1
PLACE1011054	87.07	91.74	90.14	1.05	1.04
PLACE1011056	234.93	341.05	191.57	1.45	0.82
PLACE1011057	50.32	61.94	47.85	1.23	0.95
PLACE1011059	28.75	30.18	17.75	1	1
PLACE1011066	40.74	45.85	31.71	1.13	0.98
PLACE1011087	126.9	97.7	103.08	0.77	0.81
PLACE1011090	43.14	63.67	54.04	1.48	1.25
PLACE1011109	72.28	109.31	75.78	1.51	1.05
PLACE1011114	40.38	55.61	44.13	1.38	1.09
PLACE1011116	59.89	60.46	61.74	1.01	1.03
PLACE1011122	23.85	22.65	27.32	1	1
PLACE1011133	29.8	25.5	32.54	1	1
PLACE1011134	702.54	855.96	742.75	1.22	1.06
PLACE1011143	18.05	26.08	15.76	1	1
PLACE1011146	27.97	31.76	30.52	1	1
PLACE1011160	27.65	29.84	36.02	1	1
PLACE1011165	29.12	25.53	32.58	1	1
PLACE1011181	719.05	676.22	532.05	0.94	0.74
PLACE1011185	128.61	184.54	128.11	1.43	1
PLACE1011186	161.64	160.59	193.09	0.99	1.19
PLACE1011203	26.77	26.81	25.39	1	1
PLACE1011214	745.48	767.22	632.47	1.03	0.85
PLACE1011219	38.42	58.33	37.91	1.46	1
PLACE1011221	31.09	40.42	44.12	1.01	1.1

(717)

特開 2002-191363

1431

1432

PLACE1011229	20.84	29.07	17.82	1	1
PLACE1011231	20.61	29.22	21.09	1	1
PLACE1011236	553.05	663.66	472.59	1.2	0.85
PLACE1011247	162.35	152.62	169.63	0.94	1.04
PLACE1011263	28.78	26.07	29.73	1	1
PLACE1011273	22.76	23.17	22.03	1	1
PLACE1011278	72.55	79.44	63.03	1.09	0.87
PLACE1011289	23.79	33.63	31.25	1	1
PLACE1011291	134.38	169.36	147.47	1.26	1.1
PLACE1011296	30.64	34.19	32.38	1	1
PLACE1011310	29.41	24.49	28.94	1	1
PLACE1011311	80.01	83.34	61.28	1.04	0.77
PLACE1011321	63.79	59.32	62.43	0.93	0.98
PLACE1011325	21.43	25.41	19.82	1	1
PLACE1011332	46.92	65.48	63.02	1.4	1.34
PLACE1011340	96.73	105.33	93.22	1.09	0.96
PLACE1011353	87.13	92.87	77.19	1.07	0.89
PLACE1011360	73.75	62.56	56.98	0.85	0.77
PLACE1011364	40.61	42.21	47	1.04	1.16
PLACE1011365	31.47	37.76	31.3	1	1
PLACE1011371	1393.0	1511.96	1103.92	1.09	0.79
PLACE1011375	21.17	21.56	18.08	1	1
PLACE1011386	52.19	45.65	48.93	0.87	0.94
PLACE1011399	34.74	30.13	32.85	1	1
PLACE1011406	30.27	36.26	23.82	1	1
PLACE1011407	36.4	42.05	33.38	1.05	1
PLACE1011419	27.2	44.71	42.78	1.12	1.07
PLACE1011433	37.98	29.32	35.67	1	1
PLACE1011440	18.75	14.39	23.74	1	1
PLACE1011452	52.41	64.26	64.32	1.23	1.23
PLACE1011465	25.85	23.7	25.01	1	1
PLACE1011472	25.31	25.72	28.29	1	1
PLACE1011477	589.52	635.4	447.32	1.08	0.76
PLACE1011478	115.07	140.27	106.64	1.22	0.93
PLACE1011492	72	97.27	57.14	1.35	0.79
PLACE1011498	17.1	22.26	16.32	1	1
PLACE1011501	17.79	17.26	18.01	1	1
PLACE1011503	9.79	7.51	7.6	1	1
PLACE1011509	34.49	28.69	32.22	1	1
PLACE1011514	223.19	247.57	222.01	1.11	0.99
PLACE1011516	101.6	143.7	107.63	1.41	1.06
PLACE1011520	27.41	24.9	24.32	1	1
PLACE1011538	29.14	23.25	16.78	1	1
PLACE1011555	23.6	20.53	25.6	1	1
PLACE1011561	24.88	25.71	37.82	1	1
PLACE1011563	37.3	42.35	32.36	1.06	1
PLACE1011567	59.12	55.36	61.57	0.94	1.04
PLACE1011569	110.07	98.25	92.78	0.89	0.84
PLACE1011576	107.13	119.56	82.05	1.12	0.77
PLACE1011586	58.35	60.15	53.06	1.03	0.91

(718)

特開 2002-191363

1433

1434

PLACE1011635	126.85	122.41	79.4	0.96	0.63
PLACE1011641	21.95	15.08	12	1	1
PLACE1011642	96.47	131.07	98.89	1.36	1.03
PLACE1011643	27.01	32.97	23.44	1	1
PLACE1011646	311.38	371.3	292.59	1.19	0.94
PLACE1011649	39.42	40.05	38.15	1	1
PLACE1011650	23.59	25.05	20.03	1	1
PLACE1011661	92.23	95.48	84.22	1.04	0.91
PLACE1011664	62.94	63.99	43.95	1.02	0.7
PLACE1011672	22.5	26.83	22.49	1	1
PLACE1011675	20.43	16.82	14.39	1	1
PLACE1011682	34.22	33.4	46.45	1	1.16
PLACE1011708	80.24	92.72	58.81	1.16	0.73
PLACE1011719	61.08	68.74	59.57	1.13	0.98
PLACE1011725	39.29	49.72	30.96	1.24	1
PLACE1011729	48.33	49.66	37.62	1.03	0.83
PLACE1011741	42.11	42.21	26.14	1	0.95
PLACE1011749	98.54	120.28	81.54	1.22	0.83
PLACE1011757	200.1	214.18	192.99	1.07	0.96
PLACE1011762	31.14	34.16	27.67	1	1
PLACE1011778	20.12	27.07	17.12	1	1
PLACE1011783	79.37	103.33	80.61	1.3	1.02
PLACE1011795	62.52	74.53	49.61	1.19	0.79
PLACE1011810	26.95	35.71	25.38	1	1
PLACE1011824	94.06	261.39	163.66	2.78	1.74
PLACE1011825	152.66	173.17	154.8	1.13	1.01
PLACE1011835	23.8	27.8	43.04	1	1.08
PLACE1011836	120.62	135.76	141.71	1.13	1.17
PLACE1011847	285.54	294.38	284.53	1.03	1
PLACE1011855	22.68	27.11	28.04	1	1
PLACE1011858	36.79	39.34	31.86	1	1
PLACE1011874	45.52	57.52	57.05	1.26	1.25
PLACE1011875	21.91	19.56	19.38	1	1
PLACE1011877	31.1	30.5	28.95	1	1
PLACE1011891	17.01	14.38	16	1	1
PLACE1011896	9.03	4.47	5.22	1	1
PLACE1011920	17.59	18.35	15.64	1	1
PLACE1011922	14.43	22.14	26.94	1	1
PLACE1011923	318.75	346.7	273.3	1.09	0.86
PLACE1011937	47.17	42.55	44.49	0.9	0.94
PLACE1011939	72	86.06	70.82	1.2	0.98
PLACE1011940	48.98	53.18	48.23	1.09	0.98
PLACE1011962	47.89	49.93	54.67	1.04	1.14
PLACE1011964	25.14	30.6	30.08	1	1
PLACE1011978	45.43	47.26	54.6	1.04	1.2
PLACE1011980	45.74	54.67	59.79	1.2	1.31
PLACE1011981	248.01	278.49	169.77	1.12	0.68
PLACE1011982	22.97	26.02	26.36	1	1
PLACE1011995	53.57	49.73	50.05	0.93	0.93
PLACE1012023	21.43	23.58	20.15	1	1

(719)

特開 2002-191363

1435

1436

PLACE1012026	22.39	20.44	21.38	1	1
PLACE1012031	29.21	36.05	24.28	1	1
PLACE2000003	171.38	188.33	175.86	1.1	1.03
PLACE2000005	17.08	13.58	14.3	1	1
PLACE2000006	215.06	220.3	169.37	1.02	0.79
PLACE2000007	32.91	35.43	30.39	1	1
PLACE2000011	77.03	82.84	64.01	1.08	0.83
PLACE2000014	635.17	607.14	482.26	0.96	0.76
PLACE2000015	37.29	41.56	31.59	1.04	1
PLACE2000017	14.69	19.74	13.51	1	1
PLACE2000021	44.21	63.31	49.29	1.43	1.11
PLACE2000022	46.15	49.28	51.9	1.07	1.12
PLACE2000030	1694.2	2013.81	1419.12	1.19	0.84
PLACE2000032	60.37	68.14	58.24	1.13	0.96
PLACE2000033	110.17	111.61	98.7	1.01	0.9
PLACE2000034	57.78	58.66	54.29	1.02	0.94
PLACE2000039	120.07	145.25	131.21	1.21	1.09
PLACE2000043	346.07	354.6	252.31	1.02	0.73
PLACE2000044	23.97	29.58	29.62	1	1
PLACE2000047	102.56	112.06	113.24	1.09	1.1
PLACE2000050	71.89	90.85	58.04	1.26	0.81
PLACE2000061	17.86	20.56	15.09	1	1
PLACE2000062	66.1	72.31	49.98	1.09	0.76
PLACE2000072	17.99	18.55	16.31	1	1
PLACE2000073	15.95	15.72	13.57	1	1
PLACE2000097	213.87	252.44	257.53	1.18	1.2
PLACE2000100	78.87	88.68	57.86	1.12	0.73
PLACE2000103	54.73	54.47	56.01	1	1.02
PLACE2000106	135.55	158.65	131.58	1.17	0.97
PLACE2000111	87.67	97.31	56.96	1.11	0.65
PLACE2000115	17.7	16.8	12.26	1	1
PLACE2000118	108.86	115.04	163.24	1.06	1.5
PLACE2000124	816.3	718.85	492.7	0.88	0.6
PLACE2000132	13.87	16.39	13.31	1	1
PLACE2000136	13.17	14.88	13.46	1	1
PLACE2000137	22.14	19.06	19.32	1	1
PLACE2000140	98.61	148.4	126.8	1.5	1.29
PLACE2000147	21.9	25.66	17.02	1	1
PLACE2000153	14.7	11.12	14.66	1	1
PLACE2000164	20.49	18.37	20.21	1	1
PLACE2000170	125.31	145.03	117.3	1.16	0.94
PLACE2000172	17.54	16.31	16.59	1	1
PLACE2000173	70.78	80.23	58.69	1.13	0.83
PLACE2000174	57.95	64.79	58.03	1.12	1
PLACE2000176	43.43	41.92	40.16	0.97	0.92
PLACE2000187	60.61	54.14	36.42	0.89	0.66
PLACE2000216	59.22	72.23	56.49	1.22	0.95
PLACE2000219	39.68	48.98	40.11	1.22	1
PLACE2000221	135.95	164.16	158.94	1.21	1.17
PLACE2000223	5.6	6.49	3.36	1	1

(720)

特開 2002-191363

1437

1438

PLACE2000231	33.93	38.68	26.47	1	1
PLACE2000235	98.28	112.79	92.76	1.15	0.94
PLACE2000246	94.02	98.46	83.22	1.05	0.89
PLACE2000264	79.88	92.69	64.8	1.16	0.81
PLACE2000274	35.59	37.8	39.08	1	1
PLACE2000287	20.39	20.25	23.46	1	1
PLACE2000296	18.03	15.38	14.93	1	1
PLACE2000302	45.55	58.9	46.59	1.29	1.02
PLACE2000305	155.84	169.39	135.63	1.09	0.87
PLACE2000317	32.82	20.28	20.47	1	1
PLACE2000324	41.35	38.7	36.15	0.97	0.97
PLACE2000334	183.82	175.14	175.68	0.95	0.96
PLACE2000335	136	196.92	135.48	1.45	1
PLACE2000340	16.11	18.58	11.97	1	1
PLACE2000341	210.96	277.16	209.16	1.31	0.99
PLACE2000342	166.99	114.91	107.63	0.69	0.64
PLACE2000347	154.45	189.66	153.52	1.23	0.99
PLACE2000357	79.73	96.17	102.29	1.21	1.28
PLACE2000358	256.57	305.62	211.22	1.19	0.82
PLACE2000359	3.81	4.5	1.37	1	1
PLACE2000366	99.02	132.98	84.55	1.34	0.85
PLACE2000371	68.55	85.66	71.92	1.25	1.05
PLACE2000373	43.55	47.22	46.75	1.08	1.07
PLACE2000374	23.53	32.86	28.16	1	1
PLACE2000379	16.39	17.7	16.49	1	1
PLACE2000386	1286.2	1518.95	1572.54	1.18	1.22
PLACE2000388	50.87	49.22	36.25	0.97	0.79
PLACE2000392	319.51	348.31	353.39	1.09	1.11
PLACE2000394	71.84	67.92	78.67	0.95	1.1
PLACE2000398	30.22	38.93	35.07	1	1
PLACE2000399	42.9	34.91	50.88	0.93	1.19
PLACE2000402	32.87	31.76	32.88	1	1
PLACE2000404	58.57	67.45	73.86	1.15	1.26
PLACE2000411	52.82	83.73	115.98	1.59	2.2
PLACE2000418	50.46	65.86	56.08	1.31	1.11
PLACE2000419	102.69	137.4	124.2	1.34	1.21
PLACE2000425	37.31	40.72	33.56	1.02	1
PLACE2000427	21.66	25.88	27.14	1	1
PLACE2000433	50.87	47.44	43.55	0.93	0.86
PLACE2000435	21.67	16.61	17.84	1	1
PLACE2000438	19.41	25.68	29.91	1	1
PLACE2000450	81.66	92.34	97.83	1.13	1.2
PLACE2000455	18.01	22.78	20.07	1	1
PLACE2000458	55.64	68.4	59.18	1.23	1.06
PLACE2000464	29.94	27.76	24.02	1	1
PLACE2000465	213.81	251.88	207.06	1.18	0.97
PLACE2000473	836.58	983.57	806.98	1.18	0.96
PLACE2000477	16.45	29.51	16.05	1	1
PLACE3000004	87.82	83.69	80.62	0.95	0.92
PLACE3000009	1606.2	1262.54	1148.63	0.79	0.72

(721)

特開 2002-191363

1439

1440

PLACE3000020	760.66	812.44	697.12	1.07	0.92
PLACE3000029	217.19	259.74	201.46	1.2	0.93
PLACE3000038	41.85	54.63	40.3	1.31	0.96
PLACE3000052	367.89	489.29	371.66	1.33	1.01
PLACE3000059	24.83	27.37	23.53	1	1
PLACE3000067	59.74	70.09	61.55	1.17	1.03
PLACE3000069	93.54	91.2	73.59	0.97	0.79
PLACE3000070	185.28	188.38	229.21	1.02	1.24
PLACE3000103	65.17	61.33	54.93	0.94	0.84
PLACE3000119	3.61	113.76	2.84	2.84	1
PLACE3000121	1345.4	1349.82	1036.69	1	0.77
PLACE3000124	97.43	115.22	90.24	1.18	0.93
PLACE3000135	11.35	10.57	7.29	1	1
PLACE3000136	28.24	30.03	24.93	1	1
PLACE3000142	21.84	22.96	23.32	1	1
PLACE3000145	738.49	611.83	451.51	0.83	0.61
PLACE3000147	176.32	164.52	132.27	0.93	0.75
PLACE3000148	4.21	2.12	1.47	1	1
PLACE3000154	35.29	60.89	56.38	1.52	1.41
PLACE3000155	66.71	65.47	53.88	0.98	0.81
PLACE3000156	18.22	36.57	17.58	1	1
PLACE3000157	26.5	33.22	27.53	1	1
PLACE3000158	115.37	133.81	102.19	1.16	0.89
PLACE3000160	136.75	99.83	114.37	0.73	0.84
PLACE3000169	165.8	195.53	168.21	1.18	1.01
PLACE3000181	127.79	141.83	120.13	1.11	0.94
PLACE3000194	49.19	48.32	44.38	0.98	0.9
PLACE3000197	12.91	14.38	16.78	1	1
PLACE3000199	14.71	10.56	10.36	1	1
PLACE3000205	868.43	1057.62	753.31	1.22	0.87
PLACE3000207	216.92	252.67	191.1	1.16	0.88
PLACE3000208	54.89	54.23	48.79	0.99	0.89
PLACE3000213	33.26	40.23	30.49	1.01	1
PLACE3000215	21.84	17.5	13.39	1	1
PLACE3000218	13.41	11.66	9.09	1	1
PLACE3000220	114.32	105.77	107.52	0.93	0.94
PLACE3000221	437.38	475.47	358.38	1.09	0.82
PLACE3000225	48.79	61.71	52.94	1.26	1.09
PLACE3000226	89.7	98.83	76.04	1.1	0.85
PLACE3000230	26.28	30.38	53.41	1	1.34
PLACE3000231	34.3	53.63	46.02	1.34	1.15
PLACE3000235	105.91	116.52	94.01	1.1	0.89
PLACE3000242	62.07	67.67	62.09	1.09	1
PLACE3000244	26.87	27.09	18.22	1	1
PLACE3000253	20.69	18.14	18.91	1	1
PLACE3000254	59.24	71.24	61.14	1.2	1.03
PLACE3000271	210.99	227.81	179.99	1.08	0.85
PLACE3000276	22.03	19.83	15.86	1	1
PLACE3000304	283.54	350.52	301.89	1.24	1.06
PLACE3000309	48.91	70.41	44.97	1.44	0.92

(722)

特開 2002-191363

1441

1442

PLACE3000310	42.78	41.15	34.91	0.96	0.94
PLACE3000320	49.02	51.22	40.77	1.04	0.83
PLACE3000322	3.16	0.57	1.79	1	1
PLACE3000330	221.71	298.07	276.82	1.34	1.25
PLACE3000331	138.49	160.45	123.87	1.16	0.89
PLACE3000336	37.63	52.57	39.67	1.31	1
PLACE3000339	104.65	114.64	90.45	1.1	0.86
PLACE3000341	114.82	141.48	110.97	1.23	0.97
PLACE3000350	160.43	173.42	121.49	1.08	0.76
PLACE3000352	47.6	57.89	40.6	1.22	0.85
PLACE3000353	61.18	67.39	56.69	1.1	0.93
PLACE3000362	76	90.79	76.16	1.19	1
PLACE3000363	16.4	27.93	15.48	1	1
PLACE3000365	27.86	33.26	31.95	1	1
PLACE3000373	21.61	21.8	22.19	1	1
PLACE3000374	31.32	41.25	22.3	1.03	1
PLACE3000387	8.25	9.82	6.88	1	1
PLACE3000388	64.69	77.9	44.31	1.2	0.68
PLACE3000399	112.65	133.33	109.36	1.18	0.97
PLACE3000400	44.55	49.98	41.2	1.12	0.92
PLACE3000401	490.44	600.16	510.14	1.22	1.04
PLACE3000402	32.95	41.75	27.52	1.04	1
PLACE3000405	53.52	61.97	62.72	1.16	1.17
PLACE3000406	35.93	38.71	42.23	1	1.06
PLACE3000413	18.05	16.88	18	1	1
PLACE3000416	29.64	28.6	31.16	1	1
PLACE3000425	50.28	54.42	57.61	1.08	1.15
PLACE3000437	204.43	214.99	215.82	1.05	1.06
PLACE3000455	70.16	96.18	109.86	1.37	1.57
PLACE3000475	146.58	174.44	167.87	1.19	1.15
PLACE3000477	71.98	48.83	46.22	0.68	0.64
PLACE4000003	23.25	25.51	19.95	1	1
PLACE4000008	70.24	63.65	74.26	0.91	1.06
PLACE4000009	43.12	37.54	47.82	0.93	1.11
PLACE4000014	21.64	19.96	21.69	1	1
PLACE4000029	445.92	441.3	386.76	0.99	0.87
PLACE4000034	90.28	104.98	136.62	1.16	1.51
PLACE4000049	80.02	77.88	91.08	0.97	1.14
PLACE4000052	28.33	17.89	22.57	1	1
PLACE4000062	33.61	37.15	30.93	1	1
PLACE4000063	43.96	48.75	45.39	1.11	1.03
PLACE4000089	59.97	63.79	58.77	1.06	0.98
PLACE4000093	19.92	15.59	24.78	1	1
PLACE4000100	59.02	65.05	68.87	1.1	1.17
PLACE4000103	23.19	22.13	16.6	1	1
PLACE4000106	42.56	54.08	43.9	1.27	1.03
PLACE4000128	67.58	75.38	71.3	1.12	1.06
PLACE4000129	29.93	24.81	23.87	1	1
PLACE4000131	782.86	711.57	587.61	0.91	0.75
PLACE4000147	14.59	9.44	6.98	1	1

(723)

特開 2002-191363

1443

1444

PLACE4000156	163.18	267.14	159.69	1.64	0.98
PLACE4000175	14.89	13.1	10.66	1	1
PLACE4000190	628.33	504.54	459.39	0.8	0.73
PLACE4000192	28.59	35.85	25.67	1	1
PLACE4000206	77.21	57.16	80.21	0.74	1.04
PLACE4000211	79.65	101.47	63.07	1.27	0.79
PLACE4000214	26	17.61	16.37	1	1
PLACE4000222	84.71	90.07	69.78	1.06	0.82
PLACE4000223	22.62	14.09	14.21	1	1
PLACE4000229	33.82	41.71	44.09	1.04	1.1
PLACE4000230	85.95	57.35	62.17	0.67	0.72
PLACE4000233	73.47	68.6	63.73	0.93	0.87
PLACE4000239	100.72	94.05	95.4	0.93	0.95
PLACE4000247	34.55	28.2	23.03	1	1
PLACE4000250	51.35	59.3	45.61	1.15	0.89
PLACE4000252	20.75	19.51	18.21	1	1
PLACE4000259	250.69	228.39	179.09	0.91	0.71
PLACE4000261	21.5	12.14	17.49	1	1
PLACE4000264	38.97	41.66	55.02	1.04	1.38
PLACE4000269	47.1	37.8	32.16	0.85	0.85
PLACE4000270	24.24	29.97	22.47	1	1
PLACE4000281	211.22	274.25	267.89	1.3	1.27
PLACE4000300	24.03	26.73	16.79	1	1
PLACE4000320	47.61	52.92	41.99	1.11	0.88
PLACE4000323	70.64	77.63	73.66	1.1	1.04
PLACE4000326	17.15	19.53	21.91	1	1
PLACE4000344	20.15	17.55	19.07	1	1
PLACE4000347	42.11	35.66	37.07	0.95	0.95
PLACE4000354	20.54	16.97	13.97	1	1
PLACE4000367	19.08	21.62	13.54	1	1
PLACE4000369	18.56	16.86	16.32	1	1
PLACE4000379	44.93	48.22	35.27	1.07	0.89
PLACE4000387	28.48	18.41	23.23	1	1
PLACE4000392	18.72	23.43	16.99	1	1
PLACE4000399	417.14	435.88	349.26	1.04	0.84
PLACE4000401	13.12	19.41	12.77	1	1
PLACE4000403	72.03	64.17	68.86	0.89	0.96
PLACE4000411	59.36	51.92	41.47	0.87	0.7
PLACE4000415	23.89	20.82	25.63	1	1
PLACE4000416	174.35	147.12	179.04	0.84	1.03
PLACE4000424	19.71	16.25	17.79	1	1
PLACE4000431	263.83	293	219.65	1.11	0.83
PLACE4000443	20.1	8.3	13.07	1	1
PLACE4000445	71.62	72.15	70.79	1.01	0.99
PLACE4000450	147.23	126.29	117.48	0.86	0.8
PLACE4000455	62.39	48.46	44.08	0.78	0.71
PLACE4000465	135.53	132.4	107.77	0.98	0.8
PLACE4000466	1046.6	1140.39	799.29	1.09	0.76
PLACE4000472	121.8	126.78	113.94	1.04	0.94
PLACE4000487	264.78	281.84	202.36	1.06	0.76

(724)

特開 2002-191363

1445

1446

PLACE4000489	48.45	36.77	38.35	0.83	0.83
PLACE4000494	23.84	20.8	21.4	1	1
PLACE4000502	88.64	86.75	81.9	0.98	0.92
PLACE4000521	323.04	404.56	285.14	1.25	0.88
PLACE4000522	30.08	33.41	25.49	1	1
PLACE4000537	29.74	23.24	26.85	1	1
PLACE4000548	42.55	40.31	87.2	0.95	2.05
PLACE4000558	23.58	23.82	17.37	1	1
PLACE4000581	61.42	69.56	63.92	1.13	1.04
PLACE4000590	8.23	5	6.05	1	1
PLACE4000593	18.6	17.3	14.81	1	1
PLACE4000612	41.95	53.37	32.82	1.27	0.95
PLACE4000638	26.8	37.75	22.79	1	1
PLACE4000650	27.3	16.4	33.1	1	1
PLACE4000651	36.82	39.3	38.22	1	1
PLACE4000654	16.89	15.19	15.24	1	1
PLACE4000670	16.92	13.19	15.71	1	1
PLACE4000685	187.86	211.77	202.76	1.13	1.08
PLACE4000687	16.21	22.59	13.56	1	1
PLACE5000003	21.87	28.54	21.39	1	1
PLACE5000005	110.24	120.37	147.87	1.09	1.34
PLACE5000019	13.52	8.92	10.26	1	1
PLACE5000021	13.23	15.67	10.24	1	1
PLACE5000022	19.05	19.6	21.59	1	1
PLACE5000024	34.25	21.2	21.25	1	1
PLACE5000036	40.53	31.28	35.69	0.99	0.99
PLACE5000059	352.37	319.61	357.62	0.91	1.01
PLACE5000076	47.21	58.74	32.77	1.24	0.85
PLACE5000117	84.84	102.57	87.82	1.21	1.04
PLACE5000143	23.5	22.78	23.28	1	1
PLACE5000152	15.35	12.76	11.52	1	1
PLACE5000154	144.51	199.46	237.34	1.38	1.64
PLACE5000155	89.36	81.41	79.91	0.91	0.89
PLACE5000165	78.93	76.87	88.02	0.97	1.12
SKNMC1000004	101.25	99.46	95.02	0.98	0.94
SKNMC1000011	46.1	44.65	56.16	0.97	1.22
SKNMC1000013	16.47	25.9	14.18	1	1
SKNMC1000014	92.84	113.71	75.96	1.22	0.82
SKNMC1000018	54.44	47.21	45.13	0.87	0.83
SKNMC1000020	21.7	28.63	29.3	1	1
SKNMC1000046	26.3	28.76	16.18	1	1
SKNMC1000050	62.13	60.84	53.16	0.98	0.86
SKNMC1000062	330.04	282.45	310.59	0.86	0.94
SKNMC1000075	17.55	17.79	15.1	1	1
SKNMC1000082	22.09	43.54	26.48	1.09	1
SKNMC1000091	55.45	72.12	67.23	1.3	1.21
SKNMC1000099	15.48	12.31	17.8	1	1
SKNMC1000104	19.72	17.77	17.75	1	1
SKNMC1000113	25.72	23.37	13.13	1	1
SKNMC1000119	31.7	33.07	28.84	1	1

(725)

特開 2002-191363

1447

1448

SKNMC1000142	13.17	9.46	8.15	1	1
SKNMC1000170	18.9	35.22	13.42	1	1
SKNMC1000178	36.62	37.44	34.52	1	1
SKNMC1000194	30.86	16.71	25.89	1	1
SKNMC1000198	34.11	45.61	47.99	1.14	1.2
SKNMC1000225	26.81	26.95	26.7	1	1
SKNMC1000249	8.56	7.08	5.55	1	1
SPLN1000007	23.41	31.55	22.1	1	1
SPLN1000012	12.79	13.7	8.13	1	1
SPLN1000014	70.11	67.87	59.13	0.97	0.84
SPLN1000036	594.83	558.47	477.44	0.94	0.8
SPLN1000059	18.59	12.25	11.97	1	1
SPLN1000068	70.13	64.3	54.94	0.92	0.78
SPLN1000072	32.05	40.98	29.41	1.02	1
SPLN1000101	192.61	203.71	206.97	1.06	1.07
SPLN1000108	11.49	13.35	11.51	1	1
SPLN1000113	34.46	41.39	30.71	1.03	1
SPLN1000114	45.22	48.21	35.95	1.07	0.88
SPLN1000132	34.95	36.95	42.28	1	1.06
SPLN1000135	343.33	311.82	277.27	0.91	0.81
SPLN1000136	66.82	82.96	80.88	1.24	1.21
SPLN1000141	76.01	69.18	98.16	0.91	1.29
SPLN1000164	30.6	24.91	22.66	1	1
SPLN1000166	23.21	20.04	14.17	1	1
SPLN1000175	139.17	239.48	209	1.72	1.5
SPLN1000182	14.94	11.84	11.7	1	1
SPLN1000185	55.33	55.69	40.97	1.01	0.74
THYMU1000004	71.72	74.9	83.41	1.04	1.16
THYMU1000009	94.13	77.8	90.28	0.83	0.96
THYMU1000015	228.36	266.58	260.48	1.17	1.14
THYMU1000016	85.88	95.7	65.55	1.11	0.76
THYMU1000023	20.69	15.9	18.99	1	1
THYMU1000034	24.06	21.75	16.17	1	1
THYMU1000035	16.7	12.77	9.02	1	1
THYMU1000037	19.37	14.84	13.58	1	1
THYMU1000042	79.95	82.87	68.39	1.04	0.86
THYMU1000047	89.91	103.09	89.83	1.15	1
THYMU1000080	40.1	18.75	23.34	1	1
THYMU1000094	100.33	88.03	83.95	0.88	0.84
THYMU1000109	3394.6	3204.08	2476.84	0.94	0.73
THYMU1000127	149.43	302.07	224.65	2.02	1.5
THYMU1000130	50.99	54.49	39.21	1.07	0.78
THYMU1000137	89.89	98.22	85.61	1.09	0.95
THYMU1000146	41.95	48.71	54.46	1.16	1.3
THYMU1000159	75.18	54.17	55.6	0.72	0.74
THYMU1000163	1438.2	1056.73	808.48	0.73	0.56
THYMU1000167	37.01	32.25	20.73	1	1
THYMU1000186	41.69	49.31	32.2	1.18	0.96
THYR01000017	64.39	60.8	44.03	0.94	0.68
THYR01000026	34.47	79.34	36.05	1.98	1

(726)

特開 2002-191363

1449

1450

THYRO1000034	27.76	31.8	20.76	1	1
THYRO1000035	35.4	40.29	62.06	1.01	1.55
THYRO1000036	23.43	26.13	14.41	1	1
THYRO1000040	29.81	42.36	39.84	1.06	1
THYRO1000061	37.9	51.34	31.21	1.28	1
THYRO1000067	72.96	86.57	57.57	1.19	0.79
THYRO1000070	21.53	26.8	23.38	1	1
THYRO1000072	36.97	46.63	32.18	1.17	1
THYRO1000084	44.96	56.29	56.67	1.25	1.26
THYRO1000085	49.35	66.17	67.44	1.34	1.37
THYRO1000086	11.1	15.74	10.31	1	1
THYRO1000087	8.58	16.78	5.64	1	1
THYRO1000092	52.96	62.53	59.75	1.18	1.13
THYRO1000093	8.63	7.64	6.15	1	1
THYRO1000099	37.27	32.4	28.52	1	1
THYRO1000107	24.41	21.8	26.8	1	1
THYRO1000111	21.9	17.22	25.47	1	1
THYRO1000121	20.79	17.72	18.35	1	1
THYRO1000124	8.23	6.71	9.71	1	1
THYRO1000129	6.81	8.74	7.68	1	1
THYRO1000130	37.8	41.72	38.31	1.04	1
THYRO1000132	98.28	111.98	107.16	1.14	1.09
THYRO1000134	27.46	37.47	27.29	1	1
THYRO1000144	28.27	22.09	27.89	1	1
THYRO1000155	12.36	10.14	12.44	1	1
THYRO1000156	51.42	49.18	46.2	0.96	0.9
THYRO1000163	65.44	73.1	64.89	1.12	0.99
THYRO1000173	15.15	16	17.87	1	1
THYRO1000186	210.15	248.49	187.91	1.18	0.89
THYRO1000187	77.72	82.92	67.1	1.07	0.86
THYRO1000190	75.91	65.3	68.76	0.86	0.91
THYRO1000196	10.19	6.64	5.51	1	1
THYRO1000197	74.14	96.54	82.33	1.3	1.11
THYRO1000199	18.64	18.82	15.04	1	1
THYRO1000206	29.32	53.8	59.34	1.35	1.48
THYRO1000221	53.54	70.47	61.06	1.32	1.14
THYRO1000222	38.81	43.06	43.08	1.08	1.08
THYRO1000228	45.93	36.69	32.8	0.87	0.87
THYRO1000241	97.96	115.66	91.36	1.18	0.93
THYRO1000242	39.55	36.22	38.32	1	1
THYRO1000246	34.94	38.43	32.03	1	1
THYRO1000253	62.15	52.77	55.94	0.85	0.9
THYRO1000270	25.94	22.66	21.26	1	1
THYRO1000279	10.27	4.06	5.62	1	1
THYRO1000285	91.85	96.96	103.11	1.06	1.12
THYRO1000288	44.89	35.11	51.9	0.89	1.16
THYRO1000296	51.1	53.97	49.14	1.06	0.96
THYRO1000320	52.48	54.06	54.2	1.03	1.03
THYRO1000322	33.96	29.37	30.02	1	1
THYRO1000327	29.34	27.31	29.2	1	1

(727)

特開 2002-191363

1451

1452

THYR01000343	27.82	34.09	28.73	1	1
THYR01000345	55.7	46.72	51.21	0.84	0.92
THYR01000358	9.41	7.7	6.57	1	1
THYR01000368	19.77	17.83	17.93	1	1
THYR01000375	67.03	68.23	60.2	1.02	0.9
THYR01000381	15.46	15.89	24.65	1	1
THYR01000387	36.11	33.07	29.39	1	1
THYR01000394	57.36	48.17	50.67	0.84	0.88
THYR01000395	28.97	30.24	27.58	1	1
THYR01000400	17.65	26.72	35.19	1	1
THYR01000401	24.4	25.58	18.17	1	1
THYR01000407	16.4	16.1	17.58	1	1
THYR01000420	46.85	54.92	41.53	1.17	0.89
THYR01000438	50.01	46.6	46.81	0.93	0.94
THYR01000452	42.1	54.28	49.65	1.29	1.18
THYR01000455	8.03	4.68	3.86	1	1
THYR01000471	40.24	41.92	30.96	1.04	0.99
THYR01000481	30.3	21.61	26.84	1	1
THYR01000484	61.87	67.03	50.17	1.08	0.81
THYR01000488	20.11	25.07	21.14	1	1
THYR01000501	22.62	16.34	20.31	1	1
THYR01000502	16.19	23.43	22.05	1	1
THYR01000505	13.77	12.09	10.63	1	1
THYR01000535	225.34	256.48	247.53	1.14	1.1
THYR01000556	55.39	56.81	48.63	1.03	0.88
THYR01000558	23.81	19.28	21.37	1	1
THYR01000569	74.05	68.57	73.64	0.93	0.99
THYR01000570	41.62	41.36	28.83	0.99	0.96
THYR01000572	21.39	7.23	11.72	1	1
THYR01000573	19.98	17.86	16.98	1	1
THYR01000577	14.92	4.37	7.74	1	1
THYR01000580	35.2	44.75	27.1	1.12	1
THYR01000584	51.21	39.64	46.27	0.78	0.9
THYR01000585	22.9	30.19	25.55	1	1
THYR01000596	13.06	12.04	12.16	1	1
THYR01000602	221.44	258.24	175.93	1.17	0.79
THYR01000605	22.69	14.45	17	1	1
THYR01000615	20.64	20.24	22.7	1	1
THYR01000625	55.48	53.13	39.91	0.96	0.72
THYR01000636	67.42	74.2	69.95	1.1	1.04
THYR01000637	28.99	27.15	20.45	1	1
THYR01000641	27.36	20.13	13.73	1	1
THYR01000657	25.52	29.79	32.29	1	1
THYR01000658	144.83	157.65	111.64	1.09	0.77
THYR01000662	39.96	34.34	26.55	1	1
THYR01000666	35.48	38.48	23.93	1	1
THYR01000676	55.83	49.12	51.23	0.88	0.92
THYR01000678	17.96	23.23	20.9	1	1
THYR01000684	19.12	21.46	17.53	1	1
THYR01000694	27.86	22.19	23.82	1	1

(728)

特開 2002-191363

1453

1454

THYRO1000699	89.92	90.86	76.51	1.01	0.85
THYRO1000712	184.29	270.08	170.04	1.47	0.92
THYRO1000715	143.06	353.21	194.25	2.47	1.36
THYRO1000716	33.56	44.54	28.99	1.11	1
THYRO1000717	77.22	110.47	65.42	1.43	0.85
THYRO1000723	11.31	19.35	9.86	1	1
THYRO1000734	23.48	18.16	13.64	1	1
THYRO1000748	16.72	20.76	14.58	1	1
THYRO1000755	58.84	52.18	63.38	0.89	1.08
THYRO1000756	33.49	31	34.07	1	1
THYRO1000776	50.16	42.39	50.04	0.85	1
THYRO1000777	23.87	35.27	19.82	1	1
THYRO1000779	6.32	4.93	6.1	1	1
THYRO1000782	117.27	125.31	142.84	1.07	1.22
THYRO1000783	12.02	9.07	10.19	1	1
THYRO1000786	55.04	58.8	55.91	1.07	1.02
THYRO1000787	19.96	12.46	13.7	1	1
THYRO1000792	38.21	33.39	36.86	1	1
THYRO1000793	17.33	13.84	12.2	1	1
THYRO1000795	28.83	24.46	21.51	1	1
THYRO1000796	42.19	47.03	31.27	1.11	0.95
THYRO1000798	21.24	23.63	18.2	1	1
THYRO1000800	117.04	315.89	252.61	2.7	2.16
THYRO1000805	16.57	11.21	16.09	1	1
THYRO1000815	65.07	58.64	52.6	0.9	0.81
THYRO1000829	58.28	53.39	49.78	0.92	0.85
THYRO1000835	27.36	30.54	24.1	1	1
THYRO1000843	62.82	52.11	57.79	0.83	0.92
THYRO1000846	26.19	21.19	19.35	1	1
THYRO1000852	31.05	29.41	35.84	1	1
THYRO1000855	75.51	92.14	58.07	1.22	0.77
THYRO1000865	102.88	116.21	119.95	1.13	1.17
THYRO1000866	45.28	32.65	45.38	0.88	1
THYRO1000881	89.25	87.76	93.28	0.98	1.05
THYRO1000894	19.5	18.71	15.61	1	1
THYRO1000895	24.84	14.82	13.39	1	1
THYRO1000916	86.36	104.38	80.99	1.21	0.94
THYRO1000917	1332.1	1907.82	1793.1	1.43	1.35
THYRO1000926	21.25	19.37	21.25	1	1
THYRO1000934	13.71	17.06	13.29	1	1
THYRO1000951	18.61	27.82	26.82	1	1
THYRO1000952	28.39	29.36	28.1	1	1
THYRO1000956	15.25	6.93	8.82	1	1
THYRO1000960	12.17	13.58	16.76	1	1
THYRO1000961	37.05	31.39	30.86	1	1
THYRO1000964	13.96	17.88	15.56	1	1
THYRO1000971	22.43	35.29	29.46	1	1
THYRO1000974	81.82	86.36	97.53	1.06	1.19
THYRO1000975	77.2	76.65	60.02	0.99	0.78
THYRO1000983	57.05	64.85	57.92	1.14	1.02

(729)

特開 2002-191363

1455

1456

THYRO1000984	51.78	47.85	48	0.92	0.93
THYRO1000988	57.88	51.98	58.3	0.9	1.01
THYRO1000991	14.79	9	10.42	1	1
THYRO1000999	30.62	26.66	22.67	1	1
THYRO1001003	51.62	45.23	52.45	0.88	1.02
THYRO1001015	19.88	10.58	15.95	1	1
THYRO1001016	67.86	67.9	60.96	1	0.9
THYRO1001022	19.18	19.19	16	1	1
THYRO1001031	157.9	113.8	73.39	0.72	0.46
THYRO1001033	20.62	16.08	14.67	1	1
THYRO1001062	38.12	38.16	24.23	1	1
THYRO1001063	41.75	35.16	36.06	0.96	0.96
THYRO1001071	7.28	6.77	9.2	1	1
THYRO1001080	43.21	35.05	40.67	0.93	0.94
THYRO1001093	71.48	81.87	63.33	1.15	0.89
THYRO1001100	20.18	16.25	13.76	1	1
THYRO1001102	36.88	35.13	38.38	1	1
THYRO1001104	49.54	71.39	81.87	1.44	1.65
THYRO1001109	26.9	24.31	17.57	1	1
THYRO1001113	124.88	144.83	163.96	1.16	1.31
THYRO1001120	31.33	42.01	28.69	1.05	1
THYRO1001121	28.39	26.15	21.16	1	1
THYRO1001128	56.44	105.01	71.13	1.86	1.26
THYRO1001133	129.29	141.09	109.23	1.09	0.84
THYRO1001134	6.12	48.8	3.2	1.22	1
THYRO1001142	13.49	9.24	6.46	1	1
THYRO1001173	69.76	76.75	76.64	1.1	1.1
THYRO1001175	18.21	19.94	20.86	1	1
THYRO1001177	136.34	154.56	107.81	1.13	0.79
THYRO1001189	76.97	101.83	89.67	1.32	1.16
THYRO1001194	105.93	107.92	103.06	1.02	0.97
THYRO1001204	63.61	57.71	51.46	0.91	0.81
THYRO1001205	351.28	452.06	359.35	1.29	1.02
THYRO1001213	77.39	103.68	84.02	1.34	1.09
THYRO1001224	136.69	152.12	149.46	1.11	1.09
THYRO1001237	40.47	50.92	42.87	1.26	1.06
THYRO1001242	115.84	153.83	147.74	1.33	1.28
THYRO1001258	40.94	31.04	30.95	0.98	0.98
THYRO1001262	26.89	27.89	19.46	1	1
THYRO1001266	24.21	23.01	18.49	1	1
THYRO1001271	33.6	39.06	47.64	1	1.19
THYRO1001287	618.09	675.7	490.75	1.09	0.79
THYRO1001290	8.16	5.43	5.68	1	1
THYRO1001291	59.43	91.42	60.21	1.54	1.01
THYRO1001297	36.39	49.57	53.18	1.24	1.33
THYRO1001302	16.1	15.52	13.89	1	1
THYRO1001313	12.16	11.5	7.09	1	1
THYRO1001320	67.67	100.5	55.72	1.49	0.82
THYRO1001321	47.21	52.86	37.77	1.12	0.85
THYRO1001322	24.97	27.56	20.65	1	1

(730)

特開 2002-191363

1457

1458

THYRO1001327	22.74	18.51	20.74	1	1
THYRO1001336	67.89	117.68	67.41	1.73	0.99
THYRO1001347	30.07	15.91	22.89	1	1
THYRO1001358	33.4	31.77	32.94	1	1
THYRO1001363	29.1	29.57	27.89	1	1
THYRO1001365	22.72	20.54	17.71	1	1
THYRO1001374	55.89	36.19	39.41	0.72	0.72
THYRO1001401	67.18	66.1	79.9	0.98	1.19
THYRO1001403	36.96	35.79	36.33	1	1
THYRO1001405	60.25	66.49	51.3	1.1	0.85
THYRO1001406	99.99	118.14	125.66	1.18	1.26
THYRO1001411	162.66	177.91	174.09	1.09	1.07
THYRO1001420	281.42	206.35	163.61	0.73	0.58
THYRO1001426	295.65	347.22	278.53	1.17	0.94
THYRO1001430	66.13	74.19	77.05	1.12	1.17
THYRO1001434	32.32	68.71	43.27	1.72	1.08
THYRO1001456	36.49	32.57	41.27	1	1.03
THYRO1001457	45.96	55.07	45.99	1.2	1
THYRO1001458	37.02	45.23	50.19	1.13	1.25
THYRO1001459	169.67	160.28	164.66	0.94	0.97
THYRO1001471	33.1	25.58	22.4	1	1
THYRO1001478	23.72	21.89	21.41	1	1
THYRO1001480	332.7	411.76	349.43	1.24	1.05
THYRO1001481	120.8	159.14	137.5	1.32	1.14
THYRO1001487	120.62	158.11	135.86	1.31	1.13
THYRO1001495	82.69	84.53	88.63	1.02	1.07
THYRO1001498	24.31	46.88	45.76	1.17	1.14
THYRO1001510	26.13	19.03	19.35	1	1
THYRO1001512	1869.6	2049.81	1317.06	1.1	0.7
THYRO1001519	79.4	79.3	75.14	1	0.95
THYRO1001522	89.46	101.52	59.9	1.13	0.67
THYRO1001523	71.35	72.38	74.09	1.01	1.04
THYRO1001526	240.13	365.08	318.05	1.52	1.32
THYRO1001529	90.07	80.02	57.67	0.89	0.64
THYRO1001534	50.65	49.31	81.97	0.97	1.62
THYRO1001537	166.69	165.39	144.11	0.99	0.86
THYRO1001541	177.91	167.08	139.4	0.94	0.78
THYRO1001545	51.89	36.64	42.95	0.77	0.83
THYRO1001559	145.05	146.22	132.42	1.01	0.91
THYRO1001563	183.92	184.36	182.85	1	0.99
THYRO1001570	30.95	31.51	31.27	1	1
THYRO1001573	65.95	39.85	42.27	0.61	0.64
THYRO1001584	47.65	49.27	53.94	1.03	1.13
THYRO1001593	82.04	72.74	68.9	0.89	0.84
THYRO1001595	103.98	110.51	102.49	1.06	0.99
THYRO1001596	58.19	51.42	63.84	0.88	1.1
THYRO1001602	87.72	85.88	72.86	0.98	0.83
THYRO1001605	42.84	41.69	43.2	0.97	1.01
THYRO1001608	121.51	57.35	96.56	0.47	0.79
THYRO1001617	117.84	136.24	110.76	1.16	0.94

(731)

特開2002-191363

1459

1460

THYR01001634	47.62	45.6	45.24	0.96	0.95
THYR01001637	208.17	214.42	182.7	1.03	0.88
THYR01001641	27.43	33.65	36.4	1	1
THYR01001656	59.41	54.94	78.81	0.92	1.33
THYR01001658	164.89	276.29	264.03	1.68	1.6
THYR01001661	26.03	22.14	26.37	1	1
THYR01001671	40.07	38.84	37.89	1	1
THYR01001672	15.08	14.64	16.68	1	1
THYR01001673	142.06	146.64	96.74	1.03	0.68
THYR01001677	67.29	78.89	67.23	1.17	1
THYR01001683	134.37	158.27	162.12	1.18	1.21
THYR01001700	20.47	19.37	14.84	1	1
THYR01001702	116	117.84	102.88	1.02	0.89
THYR01001703	48.06	45.1	43.24	0.94	0.9
THYR01001706	82.92	111.31	83.86	1.34	1.01
THYR01001721	34.49	36.81	37.64	1	1
THYR01001725	156.38	135.31	116.05	0.87	0.74
THYR01001730	439.5	509.26	459.36	1.16	1.05
THYR01001738	45.7	45.43	46.39	0.99	1.02
THYR01001743	28.14	23.08	20.97	1	1
THYR01001745	17.14	21.06	16.26	1	1
THYR01001746	31.83	30.99	31.95	1	1
THYR01001770	365.71	367.44	345.23	1	0.94
THYR01001772	123.83	151.39	150.86	1.22	1.22
THYR01001778	125.3	136.46	114.03	1.09	0.91
THYR01001793	118.26	92.44	95.51	0.78	0.81
THYR01001796	35.4	35.03	31.41	1	1
THYR01001800	138.35	110.54	85.07	0.8	0.61
THYR01001803	54.17	51.87	51.91	0.96	0.96
THYR01001809	68.41	58.62	37.24	0.86	0.58
THYR01001817	108.47	135.8	146.32	1.25	1.35
THYR01001819	52.11	82.26	80.67	1.58	1.55
THYR01001828	4332.9	3758.25	2993.64	0.87	0.69
THYR01001854	310.97	325.29	244.91	1.05	0.79
THYR01001895	50.43	53.8	41.92	1.07	0.83
THYR01001907	95.74	95.16	76.26	0.99	0.8
TRACH1000006	28.12	33.72	27.35	1	1
TRACH1000013	14.62	17.35	10.88	1	1
TRACH1000074	91.95	104.52	91.83	1.14	1
TRACH1000095	24.29	25.7	21.89	1	1
TRACH1000102	89.04	112.46	84.84	1.26	0.95
TRACH1000108	18.93	22.62	9.17	1	1
TRACH1000126	56.02	51.95	46.03	0.93	0.82
TRACH1000146	55.68	61.38	44.09	1.1	0.79
TRACH1000160	6.99	1.98	2.87	1	1
TRACH1000184	135.47	176.97	305.52	1.31	2.26
VESEN1000004	30.77	39.29	26.61	1	1
VESEN1000007	47.71	47.56	40.32	1	0.85
VESEN1000013	92.2	108.84	158.66	1.18	1.72
VESEN1000028	152.94	139.07	153.59	0.91	1

(732)

特開 2002-191363

1461

1462

VESEN1000059	47.54	41	36.63	0.86	0.84
VESEN1000100	43.33	51.75	59.44	1.19	1.37
VESEN1000107	36.45	39.21	41.9	1	1.05
VESEN1000117	36.31	35.07	40.25	1	1.01
VESEN1000122	47.58	33.92	57.09	0.84	1.2
VESEN1000137	54.08	51.03	54.97	0.94	1.02
VESEN1000195	59.79	53.87	60.72	0.9	1.02
VESEN1000215	20.64	18.66	18.03	1	1
VESEN1000279	182.2	194.3	161.47	1.07	0.89
VESEN1000363	130.95	132.52	120.92	1.01	0.92
VESEN1000388	47.19	34.28	39.83	0.85	0.85
VESEN1000394	45.79	54.32	54.41	1.19	1.19
VESEN1000410	33.77	24.94	18.14	1	1
VESEN1000411	40.62	43.06	55.86	1.06	1.38
VESEN1000415	44.83	38.74	34.12	0.89	0.89
VESEN1000440	77.99	76.99	81.63	0.99	1.05
VESEN1000452	55.97	59.97	56.08	1.07	1
VESEN1000539	2520.3	1514.24	2475.51	0.6	0.98
VESEN1000554	23.28	20.08	20.19	1	1
VESEN1000557	65.42	99.26	91.48	1.52	1.4
VESEN1000575	33	42.85	36.23	1.07	1
VESEN1000585	33.26	35.72	48.21	1	1.21
VESEN1000592	14.7	10.84	12.9	1	1
VESEN1000658	53.11	50.87	53.69	0.96	1.01
VESEN1000669	156.32	171.85	194.32	1.1	1.24
VESEN1000743	30.01	26.82	23.72	1	1
VESEN1000752	616.05	621.22	670.66	1.01	1.09
VESEN1000761	102.22	128.97	144.62	1.26	1.41
VESEN2000039	76.51	106.05	108.81	1.39	1.42
VESEN2000102	27.9	33.4	39.37	1	1
VESEN2000164	75.95	85.45	57.77	1.13	0.76
VESEN2000175	16.91	16.81	13.67	1	1
VESEN2000186	47.71	41.92	47.45	0.88	0.99
VESEN2000199	210.67	251.42	274.58	1.19	1.3
VESEN2000200	31.77	34.78	30.5	1	1
VESEN2000204	42.87	27.57	46.4	0.93	1.08
VESEN2000218	233.2	310.25	266.85	1.33	1.14
VESEN2000230	51.87	57.61	55.03	1.11	1.06
VESEN2000272	66.18	71.44	47.62	1.08	0.72
VESEN2000299	33.39	36.15	34.44	1	1
VESEN2000323	122.34	153.67	120.03	1.26	0.98
VESEN2000327	25.64	38.53	30.18	1	1
VESEN2000328	287.92	426.6	459.49	1.48	1.6
VESEN2000330	210.17	635.42	400.1	3.02	1.9
VESEN2000336	19.26	21	30.8	1	1
VESEN2000354	29.98	44.87	45.25	1.12	1.13
VESEN2000378	214.41	200.95	233.25	0.94	1.09
VESEN2000379	637.63	734.74	663.65	1.15	1.04
VESEN2000397	33.94	34.23	25.21	1	1
VESEN2000416	33.88	35.1	24.86	1	1

(733)

特開2002-191363

1463

1464

VESEN2000420	13.29	9.98	11.98	1	1
VESEN2000430	82.14	123.46	143.79	1.5	1.75
VESEN2000448	13.1	15.75	14.63	1	1
VESEN2000449	77.75	116.78	106.38	1.5	1.37
VESEN2000456	16.97	16.07	16.5	1	1
VESEN2000562	141.49	130.91	105.14	0.93	0.74
VESEN2000573	8.91	8.61	6.79	1	1
VESEN2000604	13.88	12.29	11.35	1	1
VESEN2000614	226.89	314.06	318.98	1.38	1.41
VESEN2000638	17.11	18.86	13.45	1	1
VESEN2000641	30.44	33.47	35.1	1	1
VESEN2000645	114.84	103.19	111.61	0.9	0.97
Y79AA1000013	23.65	28.36	17.86	1	1
Y79AA1000030	54.3	54.62	46.92	1.01	0.86
Y79AA1000033	111.64	118.12	102.03	1.06	0.91
Y79AA1000037	88.44	96.39	66.63	1.09	0.75
Y79AA1000041	30.18	31.51	34.51	1	1
Y79AA1000059	55.17	55.82	52.18	1.01	0.95
Y79AA1000065	270.08	316.25	285.13	1.17	1.06
Y79AA1000081	261.42	374.34	508.29	1.43	1.94
Y79AA1000127	107.56	107.37	68.62	1	0.64
Y79AA1000130	127.02	147.17	116.76	1.16	0.92
Y79AA1000131	6111.2	5656.63	5788.88	0.93	0.95
Y79AA1000134	52.82	69.93	53.91	1.32	1.02
Y79AA1000143	117.86	127.82	125.02	1.08	1.06
Y79AA1000144	74.35	76.59	74.72	1.03	1
Y79AA1000150	2545.9	2718.65	2054.54	1.07	0.81
Y79AA1000153	5666.0	5284.5	5846.98	0.93	1.03
Y79AA1000166	55.33	54.66	50.83	0.99	0.92
Y79AA1000179	91.15	94.27	64.88	1.03	0.71
Y79AA1000181	46.14	50.95	39.78	1.1	0.87
Y79AA1000202	588.9	770.83	565.25	1.31	0.96
Y79AA1000207	188.44	235.03	155.99	1.25	0.83
Y79AA1000214	266.68	298.79	319.39	1.12	1.2
Y79AA1000222	86.44	101.81	106.59	1.18	1.23
Y79AA1000226	82.03	161.73	169.02	1.97	2.06
Y79AA1000227	48.01	65.96	45.99	1.37	0.96
Y79AA1000230	31.14	25.61	22.03	1	1
Y79AA1000231	118.41	136.39	99.08	1.15	0.84
Y79AA1000239	151.15	168.81	146.2	1.12	0.97
Y79AA1000258	29.42	44.77	54.63	1.12	1.37
Y79AA1000268	43.67	51.54	45.57	1.18	1.04
Y79AA1000269	47.5	40.64	44.86	0.86	0.94
Y79AA1000270	130.55	110.71	113.31	0.85	0.87
Y79AA1000280	70.79	81.14	73.63	1.15	1.04
Y79AA1000285	29.49	30.14	27.66	1	1
Y79AA1000295	35.22	30.73	33.72	1	1
Y79AA1000307	18.53	20.75	18	1	1
Y79AA1000313	49.32	51.4	49.22	1.04	1
Y79AA1000314	75.77	84.06	101.41	1.11	1.34

(734)

特開 2002-191363

1465

1466

Y79AA1000328	29.85	29.66	36.26	1	1
Y79AA1000334	23.99	28.51	35.56	1	1
Y79AA1000342	327.09	366.24	329.71	1.12	1.01
Y79AA1000346	81.6	47.85	70.26	0.59	0.86
Y79AA1000347	133.77	102.63	126.89	0.77	0.95
Y79AA1000349	108.67	107.41	141.16	0.99	1.3
Y79AA1000355	162.38	124.35	158.8	0.77	0.98
Y79AA1000368	300.14	326.89	275.96	1.09	0.92
Y79AA1000388	264.81	244.65	255.31	0.92	0.96
Y79AA1000392	29.82	42.96	63.94	1.07	1.6
Y79AA1000405	76.52	90.11	72.49	1.18	0.95
Y79AA1000410	344.28	441.37	334.83	1.28	0.97
Y79AA1000420	51.5	55.34	53.93	1.07	1.05
Y79AA1000423	99.42	111.57	110.84	1.12	1.11
Y79AA1000426	48.92	57.41	47.33	1.17	0.97
Y79AA1000432	31.27	26.61	25.92	1	1
Y79AA1000453	79.54	259.3	206.07	3.26	2.59
Y79AA1000465	17.15	42	49.86	1.05	1.25
Y79AA1000469	168.58	190.42	171.96	1.13	1.02
Y79AA1000480	43.91	47.71	38.1	1.09	0.91
Y79AA1000502	92.82	78.1	86.82	0.84	0.94
Y79AA1000521	41.82	34.78	40.15	0.96	0.96
Y79AA1000534	82.06	125.38	113.59	1.53	1.38
Y79AA1000538	184.83	261.28	173.89	1.41	0.94
Y79AA1000539	354.26	344.61	222.22	0.97	0.63
Y79AA1000540	26.42	33.12	31.6	1	1
Y79AA1000560	1601.9	1282.77	1345.84	0.8	0.84
Y79AA1000574	24.19	29.5	35.63	1	1
Y79AA1000584	48.15	39.35	38.38	0.83	0.83
Y79AA1000589	2778.1	2470.28	1588.55	0.89	0.57
Y79AA1000598	29.48	26.04	21.58	1	1
Y79AA1000600	114.94	209.4	237.16	1.82	2.06
Y79AA1000609	31.84	21.28	28.15	1	1
Y79AA1000618	63.61	66.01	76.25	1.04	1.2
Y79AA1000627	62.01	61.57	49.97	0.99	0.81
Y79AA1000636	92.84	80.65	118.18	0.87	1.27
Y79AA1000649	96.75	123.64	122.08	1.28	1.26
Y79AA1000656	1766.0	1687.12	1253.96	0.96	0.71
Y79AA1000673	27.77	24.97	21.2	1	1
Y79AA1000674	1171.6	1373.94	651.44	1.17	0.56
Y79AA1000678	36.94	44.09	34.34	1.1	1
Y79AA1000682	1983.1	1699.68	1530.39	0.86	0.77
Y79AA1000683	49.85	42.48	43.56	0.85	0.87
Y79AA1000697	242.38	296.34	355.35	1.22	1.47
Y79AA1000700	43.92	66.94	58.24	1.52	1.33
Y79AA1000702	139.07	146.21	285.74	1.05	2.05
Y79AA1000704	18.66	25.29	15.37	1	1
Y79AA1000705	44.31	44.51	31.94	1	0.9
Y79AA1000717	75.03	98.74	86.2	1.32	1.15
Y79AA1000722	41.86	34.12	41.61	0.96	0.99

(735)

特開 2002-191363

1467

1468

Y79AA1000724	72.99	67.48	66.4	0.92	0.91
Y79AA1000726	33.66	36.24	29.59	1	1
Y79AA1000734	44.87	42.21	66.46	0.94	1.48
Y79AA1000748	31.18	25.23	28.06	1	1
Y79AA1000750	122.85	138.64	121.01	1.13	0.99
Y79AA1000752	32.8	31.76	33.14	1	1
Y79AA1000774	66.23	53.85	62.7	0.81	0.95
Y79AA1000776	26.59	22.27	24.72	1	1
Y79AA1000777	129.38	122.4	131.92	0.95	1.02
Y79AA1000778	51.22	36.45	32.48	0.78	0.78
Y79AA1000782	63.92	63.79	87.96	1	1.38
Y79AA1000784	52.79	64.69	85.36	1.23	1.62
Y79AA1000794	23.36	22.73	19.03	1	1
Y79AA1000800	29.75	32.72	33.2	1	1
Y79AA1000802	30.35	26.74	20.73	1	1
Y79AA1000805	19.03	22.52	16.55	1	1
Y79AA1000814	106.9	114.58	89.95	1.07	0.84
Y79AA1000823	158.18	147.37	123.66	0.93	0.78
Y79AA1000824	36.55	29.44	33.27	1	1
Y79AA1000827	72.11	68.69	52.85	0.95	0.73
Y79AA1000831	100.55	112.39	119.5	1.12	1.19
Y79AA1000833	2837.1	2936.23	2467.7	1.03	0.87
Y79AA1000850	56.14	49.87	45.58	0.89	0.81
Y79AA1000856	86.17	85.24	87.41	0.99	1.01
Y79AA1000862	21.4	15.44	14.93	1	1
Y79AA1000876	126.25	128.25	138.28	1.02	1.1
Y79AA1000888	391.22	560.15	327.4	1.43	0.84
Y79AA1000902	91.22	102.61	97.33	1.12	1.07
Y79AA1000935	63.3	75.01	65.76	1.18	1.04
Y79AA1000959	59.37	76.73	60.41	1.29	1.02
Y79AA1000962	36.15	37.57	34.42	1	1
Y79AA1000963	96.8	192.62	232.02	1.99	2.4
Y79AA1000966	519	651.66	477.36	1.26	0.92
Y79AA1000967	19.89	292.47	251.96	7.31	6.3
Y79AA1000968	80.03	76.94	75.11	0.96	0.94
Y79AA1000969	37.01	40.02	31.68	1	1
Y79AA1000976	14.88	17.75	13.35	1	1
Y79AA1000978	51.09	73.65	56.87	1.44	1.11
Y79AA1000985	236.63	273.81	258.11	1.16	1.09
Y79AA1000989	119.43	125.67	133.77	1.05	1.12
Y79AA1000991	1240.5	994.5	880.71	0.8	0.71
Y79AA1001013	1435.4	1460.4	1293.41	1.02	0.9
Y79AA1001014	59.34	57.2	61.99	0.96	1.04
Y79AA1001019	50.26	49.6	55.51	0.99	1.1
Y79AA1001020	77.34	83.75	108.42	1.08	1.4
Y79AA1001023	30.63	37.84	28.21	1	1
Y79AA1001030	54.37	72.86	67.84	1.34	1.25
Y79AA1001035	36.18	52.22	60.57	1.31	1.51
Y79AA1001041	34.02	26.39	39.75	1	1
Y79AA1001043	104.7	132.62	114.65	1.27	1.1

(736)

特開 2002-191363

1469

1470

Y79AA1001048	36.81	40.02	39.21	1	1
Y79AA1001056	83.24	70.9	78.16	0.85	0.94
Y79AA1001061	109.03	110.8	115.01	1.02	1.05
Y79AA1001062	29.05	40.37	33.85	1.01	1
Y79AA1001068	155.79	171.69	179.73	1.1	1.15
Y79AA1001073	63.42	240.36	260.57	3.79	4.11
Y79AA1001077	141.08	122.58	123.69	0.87	0.88
Y79AA1001078	89.83	99.86	113.12	1.11	1.26
Y79AA1001081	52.76	53.5	48.66	1.01	0.92
Y79AA1001088	430.32	406.89	460.33	0.95	1.07
Y79AA1001089	92.85	116.61	115.46	1.26	1.24
Y79AA1001090	67.22	69.72	59.84	1.04	0.89
Y79AA1001105	122.65	110.5	72.39	0.9	0.59
Y79AA1001142	62.73	127.62	148.35	2.03	2.36
Y79AA1001145	99.57	109.62	105.59	1.1	1.06
Y79AA1001162	61.53	55.57	54.52	0.9	0.89
Y79AA1001167	22.2	27.76	22.05	1	1
Y79AA1001176	22.58	13.4	14.3	1	1
Y79AA1001177	33.47	34.18	31.87	1	1
Y79AA1001179	172.36	288.96	209.97	1.68	1.22
Y79AA1001185	4.57	59.88	4.92	1.5	1
Y79AA1001201	75.33	82.93	99.95	1.1	1.33
Y79AA1001205	37.54	42.63	48.27	1.07	1.21
Y79AA1001211	97.72	107.77	98.62	1.1	1.01
Y79AA1001212	95.32	102.86	78.99	1.08	0.83
Y79AA1001216	650.81	671.69	747.64	1.03	1.15
Y79AA1001228	81.54	86.06	78.83	1.06	0.97
Y79AA1001233	24.91	19.52	21.84	1	1
Y79AA1001236	41.43	68.09	69.24	1.64	1.67
Y79AA1001239	66.08	63.52	79.5	0.96	1.2
Y79AA1001240	40.45	44.69	30.33	1.1	0.99
Y79AA1001255	209.03	243.4	327.16	1.16	1.57
Y79AA1001264	77.47	114.96	155.28	1.48	2
Y79AA1001272	94.67	108.27	133.39	1.14	1.41
Y79AA1001281	18.21	17.06	16.63	1	1
Y79AA1001299	84.16	104.65	104.16	1.24	1.24
Y79AA1001312	29.58	27.93	24.52	1	1
Y79AA1001319	50.23	74.51	92.58	1.48	1.84
Y79AA1001323	29.99	42.34	39.19	1.06	1
Y79AA1001328	54.08	55.7	41.63	1.03	0.77
Y79AA1001343	4712.6	5708.31	5950.25	1.21	1.26
Y79AA1001351	30.34	22.22	24.29	1	1
Y79AA1001364	57.23	72.81	54.68	1.27	0.96
Y79AA1001367	33.46	28.07	29.13	1	1
Y79AA1001384	16.59	11.98	21.5	1	1
Y79AA1001391	22.75	18.1	18.89	1	1
Y79AA1001394	107.41	106.15	85.41	0.99	0.8
Y79AA1001402	88.44	106.63	80.2	1.21	0.91
Y79AA1001410	31.55	32.32	27.63	1	1
Y79AA1001414	89.66	99.17	102.16	1.11	1.14

(737)

特開 2002-191363

1471

1472

Y79AA1001426	26.19	30.96	29.42	1	1
Y79AA1001427	944.35	1175.39	818.75	1.24	0.87
Y79AA1001430	62.12	77.68	66.7	1.25	1.07
Y79AA1001439	106.67	110.26	142.44	1.03	1.34
Y79AA1001485	46.51	42.89	31.32	0.92	0.86
Y79AA1001493	35.54	24.61	23.88	1	1
Y79AA1001511	57.74	65.35	93.44	1.13	1.62
Y79AA1001523	37.34	66.39	30.87	1.66	1
Y79AA1001530	106.49	109.86	93.6	1.03	0.88
Y79AA1001532	99.42	111.62	100.16	1.12	1.01
Y79AA1001533	64.49	53.61	44.36	0.83	0.69
Y79AA1001541	23.65	33.4	31.44	1	1
Y79AA1001548	180.15	182.1	165.86	1.01	0.92
Y79AA1001555	57.85	54.82	39.03	0.95	0.69
Y79AA1001562	159.81	149.99	152.18	0.94	0.95
Y79AA1001581	54.55	60.53	50.36	1.11	0.92
Y79AA1001585	96.52	98.24	79.13	1.02	0.82
Y79AA1001592	62.47	74.66	67.17	1.2	1.08
Y79AA1001594	6.58	7.46	4.36	1	1
Y79AA1001603	1371.8	1354.64	1311.51	0.99	0.96
Y79AA1001613	41.9	47.86	41.96	1.14	1
Y79AA1001630	21.43	17.4	21.48	1	1
Y79AA1001647	76.41	83.21	65.45	1.09	0.86
Y79AA1001664	87	149.39	85.61	1.72	0.98
Y79AA1001665	46.95	45.27	52.61	0.96	1.12
Y79AA1001679	226.19	250.68	232.31	1.11	1.03
Y79AA1001692	36.23	50.52	46.28	1.26	1.16
Y79AA1001696	14.49	20.08	15.55	1	1
Y79AA1001705	28.44	37.49	24.51	1	1
Y79AA1001711	93.7	115.28	83.78	1.23	0.89
Y79AA1001717	21.82	24.44	17.74	1	1
Y79AA1001719	56.65	50.42	51.8	0.89	0.91
Y79AA1001727	63.97	82.15	75.03	1.28	1.17
Y79AA1001750	201.52	244.27	213.32	1.21	1.06
Y79AA1001760	1525.1	1826.08	1836.66	1.2	1.2
Y79AA1001777	27.57	22.01	20.75	1	1
Y79AA1001781	14.2	10.34	11.67	1	1
Y79AA1001787	34.67	32.77	31.46	1	1
Y79AA1001793	490.12	500.64	415.02	1.02	0.85
Y79AA1001795	20.82	15.64	16.66	1	1
Y79AA1001799	63.3	90.19	82.19	1.42	1.3
Y79AA1001800	206.49	248.75	358.98	1.2	1.74
Y79AA1001801	59.81	46	44.71	0.77	0.75
Y79AA1001803	32.78	24.73	24.59	1	1
Y79AA1001805	93.54	111.29	88.98	1.19	0.95
Y79AA1001807	156.64	104.4	119.1	0.67	0.76
Y79AA1001827	54.55	36.34	49.28	0.73	0.9
Y79AA1001846	90.02	81.03	88.49	0.9	0.98
Y79AA1001848	46.02	35.28	38.62	0.87	0.87
Y79AA1001853	54.99	60.14	72.48	1.09	1.32

(738)

特開 2002-191363

1473

1474

Y79AA1001863	55.19	71.44	82.39	1.29	1.49
Y79AA1001866	73	77.51	62.79	1.06	0.86
Y79AA1001874	12.88	7.85	8.7	1	1
Y79AA1001875	92.9	99.69	99.18	1.07	1.07
Y79AA1001907	1093.8	755.57	1038.49	0.69	0.95
Y79AA1001908	28.59	18.2	21.42	1	1
Y79AA1001923	40.94	40.51	35.36	0.99	0.98
Y79AA1001927	54.84	54.43	48.17	0.99	0.88
Y79AA1001930	25.74	40.86	42.58	1.02	1.06
Y79AA1001932	97.69	98.39	84.63	1.01	0.87
Y79AA1001933	57.63	46.41	44	0.81	0.76
Y79AA1001942	37.83	42.2	33.88	1.06	1
Y79AA1001963	342.06	324.07	294.47	0.95	0.86
Y79AA1001968	133.87	148.65	174	1.11	1.3
Y79AA1001983	31.53	29.07	29.92	1	1
Y79AA1002000	38.81	38.99	47.02	1	1.18
Y79AA1002004	37.99	102.36	131.4	2.56	3.29
Y79AA1002008	90.25	65.44	54.66	0.73	0.61
Y79AA1002012	127.69	126.92	98.41	0.99	0.77
Y79AA1002017	12.63	3.46	2.78	1	1
Y79AA1002022	130.63	151.97	106.29	1.16	0.81
Y79AA1002027	31.61	32.98	29.25	1	1
Y79AA1002050	45.8	38.26	51.32	0.87	1.12
Y79AA1002058	1897.1	1240.87	921.95	0.65	0.49
Y79AA1002060	63.24	95.77	121.69	1.51	1.92
Y79AA1002062	138.95	173.15	116.65	1.25	0.84
Y79AA1002065	241.79	213.59	256.71	0.88	1.06
Y79AA1002067	69.44	85.04	88.86	1.22	1.28
Y79AA1002069	22.45	14.55	11.72	1	1
Y79AA1002070	290.21	261.7	420.51	0.9	1.45
Y79AA1002074	4784.8	4322.02	3500.57	0.9	0.73
Y79AA1002076	23.92	22.89	30.01	1	1
Y79AA1002083	32.64	25.47	18.78	1	1
Y79AA1002084	56.93	47.45	58.61	0.83	1.03
Y79AA1002086	29.98	30.26	23.7	1	1
Y79AA1002087	339.84	473.89	305.27	1.39	0.9
Y79AA1002089	63.57	68.96	66.81	1.08	1.05
Y79AA1002093	45.69	38.45	36.91	0.88	0.88
Y79AA1002101	19.61	22.84	19.2	1	1
Y79AA1002103	37.61	37.75	35.74	1	1
Y79AA1002115	48.3	60.52	68.39	1.25	1.42
Y79AA1002121	50.52	36.73	32.42	0.79	0.79
Y79AA1002125	81.65	86.76	79.2	1.06	0.97
Y79AA1002129	33.63	31.52	18.59	1	1
Y79AA1002131	20.65	14.34	20.31	1	1
Y79AA1002139	31.44	25.87	27.48	1	1
Y79AA1002144	138.16	90.77	145.45	0.66	1.05
Y79AA1002177	53.67	57.36	47.56	1.07	0.89
Y79AA1002183	168.93	154.43	165.5	0.91	0.98
Y79AA1002202	111.2	133.56	100.48	1.2	0.9

(739)

特開 2002-191363

1475

1476

Y79AA1002204	35.34	28.87	17.92	1	1
Y79AA1002206	28.51	22.35	18.61	1	1
Y79AA1002208	36.92	36.47	44.08	1	1.1
Y79AA1002209	48.12	45.04	49.5	0.94	1.03
Y79AA1002210	29.11	36.94	29.41	1	1
Y79AA1002211	57.36	63.16	52.25	1.1	0.91
Y79AA1002213	133.51	136.73	123.74	1.02	0.93
Y79AA1002215	139.31	112.01	119.56	0.8	0.86
Y79AA1002220	69.79	59.74	58.27	0.86	0.83
Y79AA1002226	91.29	74.46	87.97	0.82	0.96
Y79AA1002229	34.29	24.76	28.27	1	1
Y79AA1002234	34.52	56.44	51.95	1.41	1.3
Y79AA1002235	73.58	67.73	74.78	0.92	1.02
Y79AA1002246	50.76	51.58	35.73	1.02	0.79
Y79AA1002258	63.15	82.64	90.15	1.31	1.43
Y79AA1002279	122.18	148.88	110.84	1.22	0.91
Y79AA1002292	61.42	72.22	67.83	1.18	1.1
Y79AA1002298	39.81	43.55	30.57	1.09	1
Y79AA1002307	32.77	30.65	23.26	1	1
Y79AA1002309	18.06	24.11	13.63	1	1
Y79AA1002311	53.31	57.19	84.46	1.07	1.58
Y79AA1002334	35.05	38.07	31.31	1	1
Y79AA1002351	42.13	61.09	58.56	1.45	1.39
Y79AA1002355	48.88	42.39	40.68	0.87	0.83
Y79AA1002361	87.11	88.66	76.9	1.02	0.88
Y79AA1002365	38.75	24.26	20.53	1	1
Y79AA1002373	43.96	55.06	28.34	1.25	0.91
Y79AA1002376	3080.7	3824.05	4481.1	1.24	1.45
Y79AA1002378	73.33	93.61	68.22	1.28	0.93
Y79AA1002381	248.36	288.51	304.13	1.16	1.22
Y79AA1002388	118.82	135.82	129.37	1.14	1.09
Y79AA1002399	36.12	30.1	32.87	1	1
Y79AA1002407	57.84	42.82	52.54	0.74	0.91
Y79AA1002413	78.77	81.36	87.31	1.03	1.11
Y79AA1002416	34.3	30.2	51.99	1	1.3
Y79AA1002429	67.91	69.81	80.19	1.03	1.18
Y79AA1002431	24.66	21.16	23.98	1	1
Y79AA1002433	27.12	18.11	23.63	1	1
Y79AA1002445	78.66	54.58	73.75	0.69	0.94
Y79AA1002461	29.04	24.84	32	1	1
Y79AA1002466	882.69	904.65	782.53	1.02	0.89
Y79AA1002471	53.74	51.26	68.91	0.95	1.28
Y79AA1002472	121.95	127.4	127.11	1.04	1.04
Y79AA1002474	53.33	40.85	47.18	0.77	0.88
Y79AA1002482	103.36	111.11	116.07	1.07	1.12
Y79AA1002487	30.92	25.8	32.51	1	1
Y79AA1002490	101.4	90.92	90.54	0.9	0.89
Y79AA1002493	107.88	125.54	105.75	1.16	0.98
ZRV6C1006278	46.63	30.08	32.23	0.86	0.86

【0641】

【表170】未分化のNT2細胞、RA存在下で培養したNT2細胞、またはRA存在下で培養してさらに阻害剤を添加して培養したNT2細胞の各cDNAの発現（表中には実施例で説明の無いクローンも含まれている）

表中、NT2は無処理細胞、NT2_RAはレチノイン酸処理細胞、NT2_RA_INHIBはレチノイン酸/阻害剤処理細胞を示す *

す。また、exp.1、exp.2、およびexp.3は、それぞれn=3の測定結果を表している。t test N/Iは、無処理の細胞とレチノイン酸処理細胞との有意差検定の結果を、t test N/Iは無処理の細胞とレチノイン酸/阻害剤処理細胞との有意差検定の結果を示す。有意差検定の結果を、: P<0.05、および**: P<0.01で示した。

Clone	NT2			NT2_RA			NT2_RA_INHIB			t test N/I	t test N/I	t test N/I
	exp.1	exp.2	exp.3	exp.1	exp.2	exp.3	exp.1	exp.2	exp.3			
GAPDH(Cr2)	3.53	1.08	0.96	2.92	2.49	2.8	1.76	2.59	1.52			
β actin(Cr2)	155.4	118	99.68	148.5	110.7	101.3	114.7	105.8	151.1			
ADRG11000005	4.01	2.03	1.55	4.05	3.63	3.6	2.27	2.93	4.24			
ADRG11000007	11.08	5.73	7.92	15.42	10.6	13.87	8.99	8.17	9.15			
ADRG11000009	1.11	0.72	1.04	1.66	1.89	1.03	1.22	1.62	1.58		*	+
ADRG11000011	4.27	2.7	2.83	4.32	4.35	3.38	2.76	3.27	3.06			
ADRG11000027	1.83	0.38	0.56	0.97	0.62	0.99	0.92	1.33	1.5			
ADRG11000039	3.65	2.58	1.37	2.92	3.36	2.75	2.25	3.51	2.7			
ADRG11000069	3.25	1.83	3.28	1.86	2.53	2.83	2.01	2.89	2.7			
ADRG11000077	13.48	10.41	6.71	19.62	17.92	22.59	11.6	16.66	19.34	*	+	
ADRG11000093	5.73	2.8	4.51	7.31	5.01	4.83	3.24	6.16	7.22			
ADRG11000099	5.64	3.42	2.08	5.59	3.73	4.24	3.98	3.98	4.06			
ADRG11000136	9.97	3.52	4.19	5.77	4.73	5.86	6.61	5.16	5.49			
ADRG11000147	23.09	13.83	11.7	14.77	14.96	14.89	17.7	13.3	19.47			
ADRG11000159	6.11	2.22	3.37	5.24	2.88	4.15	2.76	2.93	3.59			
ADRG11000160	7.16	3.48	4.19	5.94	4.99	3.41	3.95	4.67	4.25			
ADRG11000171	4.84	2.99	3.23	3.52	4.19	4.37	2.55	3.88	3.45			
ADRG11000181	5.1	3.63	2.6	3.16	4.06	2.97	2.64	3.06	3.44			
BGG11000015	13.95	6.83	6.72	9.61	9.19	10.24	9.94	10.66	10.13			
BGG11000016	15.49	5.92	7.09	11.88	11.38	8.72	11.82	10.98	10.51			
BGG11000017	7.89	2.99	3.25	4.94	4.94	4.93	3.53	4.27	3.52			
BGG11000022	8.77	5.14	5.91	7.12	7.05	4.54	5.71	5.59	5.9			
BGG11000031	4.71	2.16	2.74	4.09	3.29	3.96	4.02	3.67	2.33			
BGG11000042	6.37	5.24	3.74	5.63	6.22	4.36	4.66	5.2	4.04			
BGG11000046	19.01	12.57	9.23	12.39	15.7	12.37	8.8	10.92	9.17			
BNGH41000020	859	910.1	603	164	319.2	267.4	638.2	771.6	843.4	**	-	
BNGH41000025	5.35	2.06	2.09	2.76	2.76	3.77	4.23	2.01	3.06			
BNGH41000026	16.2	7.69	7.05	9.34	11.37	9.66	10.13	7.16	10.71			
BNGH41000027	2.31	2.18	2.5	2.9	3.01	2.82	3.68	3.48	4.21	**	+	**
BNGH41000035	14.57	8.83	9.36	10.92	9.55	14.75	15.02	15.18	12.2			
BNGH41000037	10.56	7.46	6.2	8.16	9.21	6.42	3.37	5.45	4.98			
BNGH41000043	77.1	50.83	58.45	47.64	53.99	62.67	28.12	35.48	23.44	*	-	
BNGH41000048	3.5	2.19	1.91	4.28	2.87	2.4	1.63	3.01	1.78			
BNGH41000056	2.57	2.01	1	1.91	2.63	2.15	1.41	2.4	1.79			
BNGH41000067	9.84	5.84	5.53	12.49	10.26	10.25	11.74	9.68	8.33			
BNGH41000091	3.37	2.59	1.21	3.29	3.01	1.55	2.95	2.57	2.13			
BNGH41000157	10.63	5.64	6.15	8.53	9.05	7.74	6.38	6.68	5.75			
BNGH41000169	3.77	4.34	3.82	4.9	3.48	3.32	3.4	4.16	4.19			
BNGH41000181	2.47	1.59	1.84	2.93	2.1	1.8	1.7	2.66	1.59			
BNGH41000198	8.13	4.64	3.79	5.48	4.35	5.59	4.3	4.15	4.35			
BNGH41000219	9.61	3.92	4.87	4.17	5.29	5.45	5.24	7.12	7.13			
BNGH41000229	19.61	13.28	8.68	10.86	11.27	9.36	7.9	9.5	10.85			
BNGH41000237	10.9	5.47	6.45	6.65	6.97	7.79	6.36	6.25	5.44			
BNGH41000238	4.58	7	3.45	5.91	4.68	4.34	4.33	5.44	4.22			
BNGH41000243	13.85	8.69	8.48	10.19	9.71	8.97	8.23	4.87	5.54			
BNGH41000270	5.83	2.62	2.35	2.3	3.05	3.44	2.59	3.49	1.3			
BRAWH1000004	4.19	2.83	2.48	5.04	3.15	3.26	1.44	3.45	2.05			
BRAWH1000018	4.85	1.95	2.29	7.47	8.8	8.85	8.68	6.61	7.96	**	+	*
BRAWH1000021	6.52	5.06	5.87	5.09	6.94	6.44	2.89	6.23	4.28			
BRAWH1000027	11.64	8.86	7.19	8.24	10.39	11.51	5.58	7.13	8.24			
BRAWH1000029	9.58	5.15	3.52	6.01	6.72	6	5.08	5.12	5.84			
BRAWH1000040	4.6	1.89	2.14	2.92	2.71	2.7	2.92	2.5	3.01			
BRAWH1000050	11.48	4.95	5.19	9.74	7.25	8.62	8.25	8.09	8.93			
BRAWH1000051	8.18	3.93	3.19	6.15	5.72	6.02	5.01	4.25	4.44			

【0642】

【表171】

1479

1480

BRAWH100060	2.9	2.93	1.8	3.46	3.35	2.78	2.07	3.22	2.32										
BRAWH100075	2.06	1.78	1.17	2.08	2.99	2.28	1.92	2.13	2.14										
BRAWH100091	4.56	1.87	2.1	2.73	2.22	2.25	1.42	2.46	1.83										
BRAWH100094	26.53	16.26	13.57	23.37	33.3	27.71	19.86	27.26	24.74										
BRAWH100095	11.47	5.88	3.86	6.15	6.04	6.04	6.03	4.2	5.03										
BRAWH100096	7.17	5.2	3.04	5.76	6.13	4.73	6.35	5.93	7.43										
BRAWH100097	7.61	5.42	4.3	8.36	9.37	10.77	5.92	6.56	7.12	*	+								
BRAWH100100	2.35	1.26	1.29	3.27	4.09	3.18	3.47	3.17	3.82	*	+	*	+						
BRAWH100101	15.93	5.73	7.58	15.78	16.69	15.33	10.38	7.98	10.75										
BRAWH100104	1.83	1.99	1.25	3.05	2.31	2.64	0.9	2.83	2.28	*	+								
BRAWH100107	5.24	3.06	2.53	3.69	4.48	3.14	2.51	6.62	2.54										
BRAWH100110	37.02	23.89	17.91	52.01	48.45	48.78	25.83	19.88	30.82	*	+								
BRAWH100111	13.78	8.87	6.05	12.15	10.84	10.06	10.64	8.06	9.74										
BRAWH100135	11.51	6.6	6.16	7.34	6.27	6.18	7.86	3.16	9.04										
BRAWH100190	5.57	3.61	3.06	4.88	4.05	4.63	4.28	3.62	5.01										
HEMBA1000805	2.17	2.36	2.39	3.59	3.26	3.09	2.51	1.69	3.76	**	+								
HEMBA1000806	4.88	4.08	3.07	5.64	5.07	4.69	3.89	4.34	3.69										
HEMBA1000812	7.67	9.97	9.83	7.99	7.06	6.98	3.55	5.22	3.46			**	-						
HEMBA1000820	27.08	14.56	16.3	24.94	23.63	29.76	15.51	14.38	17.35										
HEMBA1000830	7.2	6.04	4.37	4.93	6.66	4.71	4.8	4.96	7.17										
HEMBA1000834	5.42	3.03	3.13	3.92	5.81	5.53	2.45	2.65	5.55										
HEMBA1000842	10.53	5.34	5.29	12.34	15.71	15.33	6.74	5.14	8.81	*	+								
HEMBA1000845	3.35	1.45	2	3.11	2.27	3.63	2.78	2.42	2.82										
HEMBA1000846	4.44	3.21	3.62	6.34	8.01	11.1	5.61	5.39	6.03	*	+	**	+						
HEMBA1000847	3.38	2.86	1.36	3.03	2.25	2.95	2.29	1.9	1.25										
HEMBA1000848	6.35	3.98	4.34	16.75	14.72	14.62	7.09	8.13	7.75	**	+	*	+						
HEMBA1000850	1.73	0.67	0.56	1.86	1.47	1.56	1.52	2.71	1.56										
HEMBA1000853	2.66	1.5	1.58	2.81	3.5	3.13	2.37	1.92	3.37	*	+								
HEMBA1000860	4.78	3.18	2.77	4.56	4.67	4.99	3.9	4.27	4.27										
HEMBA1000872	71.82	55.34	44.63	47.17	62.62	63.43	25.66	24.24	32.66			*	-						
HEMBA1000873	2.41	1.46	1.48	2.36	2.35	2.6	1.84	2.72	2.72										
HEMBA1000876	10.02	11.17	8.35	27.94	21.02	20.27	16.4	9.49	15.31	**	+								
HEMBA1000884	3.64	2.86	3.72	4.85	4.96	4.11	5.09	5.98	4.83	*	+	*	+						
HEMBA1000887	3.12	2.56	2.1	4.7	3.46	2.58	2.59	4.09	3.28										
HEMBA1000888	1.57	0.55	0.65	1.47	0.74	0.92	1.69	2.19	2.78			*	+						
HEMBA1000891	7.82	3.65	3.58	5.14	4.68	5.32	5.87	2.69	5.02										
HEMBA1000111	3.34	2.33	2.42	4.87	5.39	5.9	3.66	3.37	3.36	**	+								
HEMBA1000121	3.69	2.19	1.8	4.54	7.02	6.59	3.95	3.3	4.32	*	+								
HEMBA1000128	4.07	1.73	1.88	3.07	3.61	4.19	4.82	5.85	5.45			*	+						
HEMBA1000129	4.83	2.28	2.77	2.81	3.65	3.39	2.57	2.73	3.94										
HEMBA1000141	2.71	2.09	1.62	4.16	2.77	4.01	2.77	3.67	1.66	*	+								
HEMBA1000146	2.9	1.3	1.8	2.65	2.28	1.73	1.61	3.65	1.83										
HEMBA1000188	26.45	13.33	17.02	31.39	35.61	38.63	19.78	16.66	26.75	*	+								
HEMBA1000154	36.53	16.72	17.93	24.12	23.55	16.21	9	9.29	13.92										
HEMBA1000156	12.63	7.55	7.3	12.13	11.18	10.85	5.44	6.27	10.52										
HEMBA1000158	14.24	5.92	4.83	15.57	17.46	14.26	10.9	12.16	12.71										
HEMBA1000168	10.07	5.72	5.58	8.47	10.06	8.07	7.36	7.05	5.56										
HEMBA1000189	3.67	1.14	1.34	3.4	2.55	2.88	1.78	2.08	2.49										
HEMBA1000185	9.44	4.05	4.26	11.53	10.93	10.36	7.42	5.5	5.94	*	+								
HEMBA1000188	2.86	1.61	0.93	2.94	2.35	3.1	1.57	1.58	1.71										
HEMBA1000193	1.27	0.58	0.24	1.37	0.89	0.82	0.26	0.53	0.45										
HEMBA1000194	11.09	4.55	5.41	17.15	17.6	13.81	11.08	8.03	17.29	*	+								
HEMBA1000201	3.51	1.9	1.75	4.07	2.62	2.46	2.06	2.69	2.83										
HEMBA1000213	2.2	0.91	0.97	1.85	2.66	1.89	1.72	1.64	1.67										
HEMBA1000216	4.38	3.53	3.49	7.1	6.02	3.1	3.46	3.84	4.14										
HEMBA1000227	6.93	1.95	2.95	5.27	3.71	3.99	3.84	2.55	3.65										

2/180

【0643】

【表172】

1483

1484

HEMBA1000481	20.13	11.47	12.73	18.55	18.55	15.53	7.84	7.33	12.91										
HEMBA1000482	7.66	4.44	4.62	7.86	6.19	6.89	3.5	5.38	6.42										
HEMBA1000490	4.18	2.68	1.34	3.92	5.37	3.63	2.12	2.88	4.31										
HEMBA1000491	7.15	3.43	2.52	5.3	6.82	6.64	4.25	3.29	3.33										
HEMBA1000498	10.26	6.11	4.98	10.58	18.06	18.44	9.53	6.44	8.57	*	+								
HEMBA1000501	10.31	9.16	7.08	7.41	5.02	8.46	4.06	4.46	3.72										
HEMBA1000504	0.29	1.06	0.88	2.55	1.79	2.74	3.2	4.91	2.54	*	+	*							
HEMBA1000505	4	3.11	2.61	4.34	3.87	4.06	3.11	3.95	3.94										
HEMBA1000507	8.99	4.59	6.64	9.35	10.47	8.63	5.53	8.59	7.24										
HEMBA1000508	8.59	6.68	6.07	11.49	13.9	16.57	7.32	8.75	9.79	*	+								
HEMBA1000510	2.98	1.78	1.55	2.04	2.31	1.71	2.15	1.54	1.87										
HEMBA1000519	13.74	9.63	6.41	18.15	26.1	23.45	14.61	12.39	16.75	*	+								
HEMBA1000520	0.74	1.54	1.42	0.53	4.99	5.32	0.3	3.24	3.21										
HEMBA1000523	2.58	1.73	1.85	2.49	2.81	3.42	2.38	3.31	2.63										
HEMBA1000531	5.39	5.46	3.11	3.93	6.67	3.26	3.72	3.54	2.94										
HEMBA1000534	0.79	3.21	2.91	1.73	9.74	6.64	0.85	6.6	3.17										
HEMBA1000538	-0.07	2.6	2.6	0.69	6.28	5.42	0.12	7.11	5.18										
HEMBA1000540	3.94	2.64	3.3	8.03	7.49	8.11	2.04	3.68	2.54	**	+								
HEMBA1000542	5.67	3.4	2.44	3.85	3.3	5.44	3.98	3.82	4.97										
HEMBA1000545	2.41	1.53	0.38	4.15	3.69	3.21	1.98	2.16	2.09	*	+								
HEMBA1000547	1.74	1.59	1.68	5.72	8.77	7.03	3.43	3.74	3.3	**	+	**	+						
HEMBA1000551	9.65	6.1	8.03	14.99	17.46	18.61	8.56	8.89	9.19	**	+								
HEMBA1000555	5.3	2	2.07	3.79	6.18	4.25	2.7	2.98	2.37										
HEMBA1000557	4.48	2.92	3.57	7.15	7.8	8.32	4.31	6.14	5.01	**	+								
HEMBA1000561	3.7	1.44	1.77	4.14	3.06	3.15	3.47	4.41	2.34										
HEMBA1000563	1.24	0.37	0.85	2.27	1.82	2.27	0.66	2.98	0.86	*	+								
HEMBA1000567	3.87	1.04	1.51	8.01	8.19	8.67	2.66	3.73	4	**	+								
HEMBA1000568	3.88	2.11	2.05	5.69	5.23	5.4	1.77	2.82	3.91	*	+								
HEMBA1000569	4.97	2.5	2.71	6.85	4.01	5.8	3.46	3.51	4.29										
HEMBA1000575	13.92	7.22	8.43	20.52	24.59	18.68	11.63	11.79	11.04	*	+								
HEMBA1000588	1.28	0.91	1.2	2.91	2.49	2.9	1.78	2.48	2.62	**	+	*	+						
HEMBA1000590	3.14	1.5	1.84	3.09	1.65	1.71	1.44	1.82	1.81										
HEMBA1000591	6.68	3.99	4.87	8.78	6.73	9.08	5.54	5.94	6.27										
HEMBA1000592	1.77	1	1.66	2.61	3.4	2.25	1.98	2.18	1.99	*	+								
HEMBA1000594	3.25	0.68	1.19	1.74	3.07	2.12	1.39	1.15	1.72										
HEMBA1000604	5.99	4.47	2.05	8.88	9.05	6.96	6.29	5.91	6.23	*	+								
HEMBA1000607	4.99	3.1	3.35	6.44	6.82	5.81	3.43	4.28	4.42	*	+								
HEMBA1000608	0.99	1.94	0.42	3.85	2.15	1.46	2.61	2.1	3.4										
HEMBA1000622	2.66	1.16	0.99	4.04	3.67	4.04	2.76	3.15	3.26	*	+								
HEMBA1000634	28.82	15.23	16.08	35.62	36.93	32.2	24.35	21.77	26.76	*	+								
HEMBA1000636	10.44	4.41	5.46	7.42	7.72	8.09	6.42	4.97	5.75										
HEMBA1000637	5.28	3.33	4.09	4.63	6.26	5.53	4.14	4.87	4.43										
HEMBA1000635	7.39	4.24	2.84	8.57	9.07	9.85	5.75	6.56	6.78	*	+								
HEMBA1000657	7.14	3.73	3.78	6.89	5.66	6.19	7.09	4.53	7.57										
HEMBA1000662	2.8	1.64	1.1	1.89	1.7	1.33	1.86	1.9	1.81										
HEMBA1000664	2.6	2.45	0.17	3.74	3.57	2.7	2.86	2.52	2.77										
HEMBA1000671	3.69	2.81	2.74	7.05	5.05	5.15	3.14	2.82	3.51	*	+								
HEMBA1000673	5.96	2.79	3.34	9.32	7.79	7.67	4.47	3.8	5.32	*	+								
HEMBA1000675	2.45	2.8	0.77	6.63	4.04	4.43	3.65	3.8	3.87	*	+	*	+						
HEMBA1000678	7.03	5.09	6.34	10.12	8.74	9.2	2.93	5.72	5.28	*	+								
HEMBA1000682	5.22	2.07	2.75	12.42	15.95	13.04	14.17	11.88	14.92	**	+	**	+						
HEMBA1000686	5.1	3.46	2.35	5.21	4.74	3.32	3.54	2.67	2.25										
HEMBA1000702	9.79	6.15	6.42	10.8	11.22	8.35	8.93	8.45	8.8										
HEMBA1000705	1.79	1.26	0.4	2.12	2.25	1.15	1.75	1.57	2.17										
HEMBA1000713	5.63	3.58	2.89	6.69	5.36	6.21	7.06	6.72	4.64										
HEMBA1000718	4.7	2.67	2.33	5.7	6	5.76	3.69	3.85	2.59	*	+								

4/180

【0645】

【表174】

1485

1486

HEMBA1000719	4.82	2.97	2.79	3.61	4.58	3.67	3.75	2.77	3.67										
HEMBA1000723	2.03	0.86	1.42	1.98	2.82	1.59	1.34	3.92	2.07										
HEMBA1000726	10.3	9.3	7.72	23.56	26.89	19.83	12.69	13.58	11.3	**	+	*	+						
HEMBA1000737	6.04	3.96	3.25	8.14	10.96	7.59	6.32	6.82	2.98	*	+								
HEMBA1000739	3.01	2.28	1.42	2.14	1.87	1.92	2.98	2.21	2.48										
HEMBA1000736	4.72	2.16	2	3.64	1.97	1.99	2.73	2.2	2.64										
HEMBA1000743	0.32	1.05	0.53	1.51	2.41	0.98	0.72	1.22	1.24										
HEMBA1000745	1.74	1.73	1.32	1.18	1.69	2.12	1.96	2.53	1.18										
HEMBA1000747	4.19	1.78	1.08	3.03	2.21	1.78	1.85	3.32	2.09										
HEMBA1000748	2.17	1.28	2.24	2.2	3.52	2.79	1.4	2.38	1.72										
HEMBA1000749	4.95	3.09	2.17	6.45	8.33	7.14	3.25	4.29	3.58	*	+								
HEMBA1000753	4.81	3.6	2.79	5.03	6.01	4.99	3.34	3.06	3.28										
HEMBA1000753	9.91	6.17	6.18	9.28	11.1	8.29	5.77	5.12	5.5										
HEMBA1000757	7.1	7.74	5.44	11.01	14.04	12.37	5.38	4.46	4.75	**	+								
HEMBA1000760	16.78	13.36	13.64	8.72	12.16	6.16	8.22	7.22	7.97										
HEMBA1000769	7.05	2.51	3.23	9	8.67	9.72	4.24	4.83	3.98	*	+								
HEMBA1000773	1.32	0.68	0.25	0.36	1.46	1.1	0.81	1.64	0.68										
HEMBA1000774	8	3.27	7.05	12.39	12.55	13.92	7.51	8.12	7.46	*	+								
HEMBA1000780	2.14	1.77	0.74	2.61	2.17	1.75	1.28	2.13	1.21										
HEMBA1000783	1.08	1.96	1.07	2.21	1.08	2.2	1.9	1.74	1.44										
HEMBA1000791	3.14	3.15	3.13	6.58	7.53	5.76	3.73	3.72	6.22	**	+								
HEMBA1000793	9.3	4	3.98	5.49	6.93	5.86	5.38	4.76	5.7										
HEMBA1000802	3.76	2.25	1.22	2.43	3.6	2.62	0.88	2.18	1.88										
HEMBA1000813	9.81	3.16	4.27	6.99	7.53	7.12	3.67	6.02	6.65										
HEMBA1000817	2.66	1.43	0.92	2.74	3.08	2.72	1.26	2.52	1.67										
HEMBA1000822	0.99	1.09	0.85	1.62	3.22	2.71	1.22	1.82	0.71	*	+								
HEMBA1000837	7.7	6.4	3.84	6.01	6.66	6.53	3.91	3.03	4.64										
HEMBA1000835	5.1	2.66	2.23	8.93	7.69	7.93	7.69	5.86	6.86	**	+	*	+						
HEMBA1000835	5.71	3.29	3.29	5.75	3.34	4.85	2.51	3.39	3.41										
HEMBA1000843	6.36	5.57	5.21	6.61	9.85	9.29	4.9	5.64	10.02										
HEMBA1000851	4.2	1.79	2.1	3.58	3.85	2.86	2.91	1.96	2.78										
HEMBA1000852	5.4	3.22	2.28	5.81	4.07	5.82	2.77	3.99	3.71										
HEMBA1000867	1.61	2.47	1.06	2.17	3.19	2.37	0.68	2.24	0.83										
HEMBA1000869	1.82	1.11	0.72	0.98	2.58	1.99	0.79	2.22	0.83										
HEMBA1000870	6.82	3.33	3.67	6.25	6.67	4.52	3.47	4.37	5.69										
HEMBA1000872	4.12	2.25	3.08	4.7	5.64	4.68	3.33	3.29	4.33	*	+								
HEMBA1000875	1.77	1.41	1.93	5.81	7.31	5.85	7.19	6.68	8.14	**	+	**	+						
HEMBA1000876	5.86	4.79	3.07	7.1	7.28	6.57	4.55	4.52	6.23	*	+								
HEMBA1000907	2.12	2.01	0.66	2.54	2.27	2.12	2.3	1.55	1.2										
HEMBA1000908	4.73	8.03	3.2	3.97	8	4.77	4.32	3.17	3.88										
HEMBA1000910	4.06	2.39	3.23	5.88	8	5.6	3.31	3.17	3.05	*	+								
HEMBA1000918	3.62	1.79	2.38	3.54	2.97	3.56	2.53	2.34	2.18										
HEMBA1000919	6.44	3.37	2.05	4.74	4.83	4.38	3.75	4.79	3.18										
HEMBA1000934	8.7	4.01	3.93	4.96	5.39	5.6	4.1	3.51	4.76										
HEMBA1000935	2.09	1.32	1.09	2.05	2.33	2	1.1	2.66	1.72										
HEMBA1000940	4.94	2.14	2.53	3.07	4.88	4.53	2.3	2.63	3.13										
HEMBA1000942	6.3	3.89	2.49	7.11	9.28	8.54	5.01	6.29	5.19	*	+								
HEMBA1000943	1.78	1.53	1.49	3.18	2.76	2.23	1.98	2.58	2.05	*	+	*	+						
HEMBA1000946	8.15	7.73	6	5.16	5.33	4.88	2.92	2.97	2.91	*	-	**	-						
HEMBA1000960	9.59	5.75	7.08	15.65	18.02	18.53	8.6	9.11	11.03	**	+								
HEMBA1000963	6.47	2.77	4.32	4.75	7	7.13	5.16	3.51	7.23										
HEMBA1000968	7	1.7	1.54	3	4.17	3.31	2.23	2.96	3.14										
HEMBA1000971	5.14	1.71	2.36	4.85	4.32	4.5	3.47	4.15	3.21										
HEMBA1000972	3.69	1.13	1.73	5.98	4.9	5.9	2.76	4.55	2.35	*	+								
HEMBA1000974	1.6	0.93	0.68	2.29	1.66	2.44	2.01	3.61	2.23	*	+								
HEMBA1000975	3.28	2	1.5	5.97	3.13	2.57	2.25	4.14	2.32										

5/180

【0646】

【表175】

1487

1488

HEMBIA1000979	5.49	2.18	2.97	6.7	3.77	4.39	3.48	5.27	4.03										
HEMBIA1000981	9.63	9.63	8.99	5.49	6.85	5.43	3.2	5.8	4.89	**	-	**	-						
HEMBIA1000983	6.43	3.92	2.91	5.46	7.35	6.51	4.3	3.18	4.68										
HEMBIA1000985	1.63	1.32	0.83	1.53	0.96	1.83	1.43	0.82	1.18										
HEMBIA1000986	8.66	3.35	3.49	7.79	10.67	12.32	6.59	5.63	7.52										
HEMBIA1000991	3.99	3.51	3.27	7.03	8.03	8.59	3.11	5.46	4.41	**	+								
HEMBIA1001007	6.98	3.16	4.1	4.53	6.32	6.25	5.08	5.14	4.03										
HEMBIA1001008	3.18	2.08	1.67	6.05	4.43	4.59	2.99	3.85	3.36		+								
HEMBIA1001009	3.19	2.06	1.89	3	2.73	3.35	2.83	4.13	2.55										
HEMBIA1001014	5.39	3.12	5.74	9.86	11.08	12.45	6.45	7.98	7.75	**	+								
HEMBIA1001017	7.4	4.83	4.74	5.73	6.28	5.4	4.08	4.41	5.88										
HEMBIA1001019	2.85	2.29	1.26	2.91	2.72	2.07	1.51	2.11	2.14										
HEMBIA1001020	3.1	1.76	1.25	4.02	4.91	3.89	2.56	2.42	2.65		+								
HEMBIA1001021	5.67	3.26	3.56	5.27	3.84	4.59	5.11	3.82	6.55										
HEMBIA1001022	4.52	3.09	3.23	5.25	4.72	3.27	2.64	3.83	3.82										
HEMBIA1001024	1.94	0.42	0.87	1.28	1.11	2.19	1.54	1.4	1.02										
HEMBIA1001026	1.87	1.27	0.7	1.76	2.89	2.28	1.38	1.06	1.68										
HEMBIA1001043	2.16	1.91	1.95	3.51	4.01	3.96	1.57	1.82	0.63	**	+								
HEMBIA1001051	12.22	4.78	5.28	19.03	15.88	16.62	10.42	7.53	10.73	**	+								
HEMBIA1001052	1.62	0.97	1.98	2.35	4.21	2.8	2.24	1.49	2.61										
HEMBIA1001059	6.89	2.24	2.49	4.96	3.77	4.83	4.31	1.48	4.43										
HEMBIA1001060	7.98	3.88	4.72	10.32	9.35	8.51	6.1	5.55	6.56	**	+								
HEMBIA1001064	5.36	3.84	3.22	6.43	5.66	4.77	2.55	3.39	3.71										
HEMBIA1001074	1.62	1.41	0.32	1.6	17.18	11.61	12.79	12.04	12.64	**	+	**	+						
HEMBIA1001077	4.45	3.8	1.96	11.6	9.35	8.57	3.08	5.61	3.95	**	+								
HEMBIA1001078	14.1	8.18	8.99	5.43	6.25	7.02	4.32	6.96	5.16										
HEMBIA1001080	7.99	3.95	2.49	3.69	5.23	5.89	3.35	4.03	3.93										
HEMBIA1001084	5.31	2.86	2.62	7.71	7.07	6.47	5.73	4.4	5.39	*	+								
HEMBIA1001085	13.38	7.46	10.01	19.29	18.48	14.18	11.36	11.18	10.99	*	+								

1489

1490

HEMBA1001231	2.19	1.18	0.78	2.28	2.36	2.53	2.4	3.63	1.51										
HEMBA1001232	1.21	1.77	1.22	2.62	2.13	1.37	0.82	1.74	2.82										
HEMBA1001236	13.52	10.49	8.9	18.36	20	19.62	7.7	10.44	7.45	**	+								
HEMBA1001238	13.05	5.12	4.29	9.55	8.22	7.69	6.04	7.48	7.86										
HEMBA1001239	12.71	9.28	6.69	8.25	7.48	7.38	10.2	8.81	12.42										
HEMBA1001235	4.86	4.97	4.74	7.89	8.06	6.71	5.12	7.06	11.33	**	+								
HEMBA1001234	5.14	3.54	3.32	7.04	6.92	8.57	3.98	4.55	5.25	*	+								
HEMBA1001242	9.9	9.56	8.33	13.88	6.68	13.26	5.82	6.16	5.11			**	-						
HEMBA1001247	4.46	1.61	1.9	3.57	3.49	3.72	3.78	3.48	3.42										
HEMBA1001253	5.27	3.3	2.61	4.73	4.85	2.62	2.61	2.92	2.88										
HEMBA1001297	3.88	2.26	2.32	3.08	5.15	4.69	1.41	2.58	1.9										
HEMBA1001261	30.79	16.66	18.37	18.07	18.08	21.82	20.19	23.46	27.67										
HEMBA1001262	2.76	4.04	1.52	6.54	5.42	3.57	2.84	3.16	4.61										
HEMBA1001265	5.5	6.7	4.27	9.23	8.19	10.09	4.34	5.27	5.82	*	+								
HEMBA1001266	7.76	6.62	6.38	9.80	9.6	8.87	6.28	5.38	7.63	**	+								
HEMBA1001269	37.26	20.56	22.9	18.88	18.77	19.35	8.45	11.29	14.06			*	-						
HEMBA1001272	1.9	1.41	1.17	1.81	2.19	2.98	1.62	1.83	1.14										
HEMBA1001279	7.18	4.53	5.66	6.03	6.98	6.47	3.39	5.47	3.9										
HEMBA1001281	5.42	5.55	6.33	11.93	16.02	13.78	5.82	4.84	7.89	**	+								
HEMBA1001286	25.93	14.58	10.17	19.52	21.77	19.41	15.05	12.01	17.84										
HEMBA1001289	4.9	3.9	2.72	4.42	4.59	5.54	4.24	2.99	5.3										
HEMBA1001291	12.14	5.79	5.07	8.25	5.62	6.51	5.37	5.12	8.98										
HEMBA1001294	3.24	2.44	2.03	4.94	4.48	4.82	2.73	2.45	3.08	**	+								
HEMBA1001296	3.68	1.57	1.28	2.91	2.24	3.02	2.56	2.34	2.65										
HEMBA1001297	5.4	4.74	4.72	5.79	6.42	4.8	3.21	2.6	2.27			**	-						
HEMBA1001299	6.03	3.81	4.28	7.69	11.74	10.72	5.99	5.39	5.03	*	+								
HEMBA1001302	6.53	3.1	5.55	4.99	5.75	7.13	4.2	5.14	4.56										
HEMBA1001303	3.57	2.21	0.92	2.41	4.91	3.42	1.52	2.66	2.14										
HEMBA1001306	22.18	12.36	12.24	18.89	23.21	22.17	16.22	12.41	17.9										
HEMBA1001308	11.41	6.87	7.33	12.58	12.35	13.73	8.36	8.24	9.57	*	+								
HEMBA1001310	7.21	5.67	6.18	9.02	7.1	8.4	7.65	6.89	8.59										
HEMBA1001312	6.83	4.78	4.59	4.91	5.69	6.9	6.83	6.24	6.66										
HEMBA1001319	9.37	9.17	0.45	0.79	0.92	1.12	0.66	2.44	0.75	**	+								
HEMBA1001322	7.21	5.19	6.74	8.06	10.08	9.08	6.21	7.42	7.75	*	+								
HEMBA1001323	4.23	3.25	2.82	10.32	10.14	7.03	8.56	8.82	9.24	**	+	**	+						
HEMBA1001326	5.74	3.25	2.25	3.17	5.59	5.42	5.13	3.49	5.64										
HEMBA1001327	2.36	2.51	1.03	2	2.41	3.09	2.74	2.46	3.87										
HEMBA1001330	5.82	3.46	4.35	11.86	14.54	13.29	6.08	7.36	9.3	**	+								
HEMBA1001348	1.13	2.19	2.78	4.2	2.23	2.88	1.71	2.63	2.74										
HEMBA1001350	12.56	10.68	7.51	15.66	13.69	14.52	11.25	9.44	10.45	*	+								
HEMBA1001351	8.18	6.48	5.91	13	14.47	12.39	10.67	8.35	8.14	**	+								
HEMBA1001352	7.26	6.11	6.06	7.73	6.7	6.17	5.26	7.29	6.09										
HEMBA1001353	31.3	26.87	27.53	25.75	22.73	20.82	12.94	17.74	19.72	*	-	**	-						
HEMBA1001358	34.05	17.05	14.31	20.81	35.28	26	9.32	9.14	12.44										
HEMBA1001361	1.82	1.14	2.1	2.53	3.2	3.65	3	2.26	2.92	*	+	*	+						
HEMBA1001364	1.53	0.54	0.65	1.45	1.91	1.58	2.49	1.92	1.51										
HEMBA1001375	3.85	2.39	2.36	2.46	4.27	4.44	3.43	3.13	3.87										
HEMBA1001377	8.53	6.83	5.73	14.04	14.14	13.21	6.15	8.25	6.71	**	+								
HEMBA1001383	2.54	1.25	1.73	3.25	2.57	3.3	1.34	2.99	1.96										
HEMBA1001387	4.07	1.84	3.25	5.31	4.33	4.27	3.43	4.91	3.24										
HEMBA1001388	4.68	4.67	4.87	6.78	7.58	7.44	4.52	4.77	5.63	**	+								
HEMBA1001390	7.44	5.12	5.37	11.6	11.73	11.98	10.29	9.6	11.06	**	+	**	+						
HEMBA1001391	1.33	1.22	1.11	3.9	2.02	2.65	2.4	1.36	1.99	*	+								
HEMBA1001398	5.47	2.84	3.19	6.23	6.58	7.24	3.41	4.42	5.15	*	+								
HEMBA1001405	5.26	1.42	2.09	2.65	2.11	2.81	3.1	2.12	3.28										
HEMBA1001406	3.16	2.03	2.11	4.74	3.74	4.54	2.55	3.04	2.7	*	+								

7/180

【0648】

【表177】

1491

1492

HEMBA1001407	5.43	1.63	2.98	3.95	4.01	3.47	2.95	2.92	2.93										
HEMBA1001411	2.17	0.69	0.63	2.51	1.83	3.63	1.29	1.35	1.63										
HEMBA1001413	5.49	2.49	2.2	4.28	3.2	3.97	3.24	2.49	2.68										
HEMBA1001414	3.79	2.32	2.38	3.06	1.8	2.44	2.65	3.53	3.21										
HEMBA1001415	6.49	2.16	2.76	5.46	6.84	6.46	4.32	4.17	5.11										
HEMBA1001416	6.22	3.74	3.23	8.62	6.54	6.82	5.91	4.3	6.02										
HEMBA1001433	5.37	2.98	3.43	7.69	6.86	7.06	3.39	4.18	4.43	*	+								
HEMBA1001435	4.8	2.47	2.21	6.26	5.3	4.79	3.29	2.49	2.37										
HEMBA1001435	8.18	4.71	5.41	14.34	11.54	13.2	6.78	6.29	7.26	**	+								
HEMBA1001443	1.65	1.46	0.73	2.67	3.31	2.57	0.77	1.88	2.03	*	+								
HEMBA1001446	9.08	2.53	3.23	6.88	6.71	6.42	5.95	6.22	6.04										
HEMBA1001450	7.08	5.32	4.43	8.06	5.46	8.96	5.99	5.4	5.68										
HEMBA1001454	10.16	4.17	5.03	16.08	14.78	15.21	9.95	9.22	10.42	**	+								
HEMBA1001455	1.25	1.28	0.63	2.33	2.23	1.74	2.53	2.34	2.01	*	**	+							
HEMBA1001459	3.35	1.42	1.26	1.83	2.02	1.94	1.14	1.39	2.31										
HEMBA1001461	8.81	3.16	4.05	10.82	10.26	6.95	6	5.33	4.95										
HEMBA1001462	2.66	2.42	2.15	2.1	1.78	2.07	1.34	1.53	2.31										
HEMBA1001463	7.17	2.73	3.52	7.24	7.08	8.95	4.33	5.14	4.39										
HEMBA1001469	7.79	8.03	2.81	8.15	8.71	7.67	5.88	4.2	6.47										
HEMBA1001473	2.06	0.9	0.31	1.64	1.59	1.3	1.54	1.11	1.32										
HEMBA1001477	1.25	0.8	0.62	0.91	1.28	0.76	1.34	2.38	1.44										
HEMBA1001478	2.09	0.93	1.34	1.5	1.78	0.98	1.62	2.3	1.59										
HEMBA1001480	12.07	6.47	7.53	8.82	7.12	9.89	6	6.87	5.33										
HEMBA1001483	4.46	3.27	2.35	2.86	3.34	4.48	1.86	2.27	1.82										
HEMBA1001490	1.81	1.4	1.03	1.82	1.46	1.52	1.48	2.37	1.32										
HEMBA1001495	36.22	21.61	21.87	15.42	21.1	17.04	16.21	19.62	20.73										
HEMBA1001497	7.26	3.96	4.28	11.8	9.61	9.83	5.21	4.28	5.2	*	+								
HEMBA1001510	13.72	5.93	6.56	13.7	15.62	12.58	10.78	9.6	9.58										
HEMBA1001515	2.6	2	0.87	2.75	3.2	2.93	2.35	3.19	2.52										
HEMBA1001517	1.89	1.95	1.22	2.95	2.33	2.76	1.72	1.66	2.42	*	+								
HEMBA1001522	3.61	1.7	1.12	1.99	2.84	1.73	1.04	1.87	1.3										
HEMBA1001526	5.16	2.43	3.68	6.63	4.1	5.88	3.55	3.16	3.42										
HEMBA1001533	8.95	4.93	4.41	7.97	8.75	10.67	4.59	5.06	4.92										
HEMBA1001547	35.19	25.44	22.4	15.45	14.19	13.27	6.7	4.99	4.47	*	**	+							
HEMBA1001552	8.07	6.24	3.86	9.62	10.94	7.97	8.18	5.74	5.97										
HEMBA1001553	16.17	10.48	11.7	14.97	19.64	15.26	19.38	22.7	26.62	*	+								
HEMBA1001557	8.77	5.74	4.35	8.02	8.99	7.7	7.33	5.59	10.39										
HEMBA1001563	3.9	1.92	1.89	5.08	3.9	4.71	2.33	3.96	2.78										
HEMBA1001566	3.98	2.49	2.79	5.22	9.83	5.76	3.59	4.31	4.01										
HEMBA1001569	8.8	4.36	5.19	13.14	14.49	14.76	6.66	7.84	10.58	**	+								
HEMBA1001578	10.01	5.49	7.22	16.18	15.76	21.41	6.88	8.18	7.08	*	+								
HEMBA1001579	24.95	9.44	8.88	11.45	10.82	11.3	6.85	6.64	9.61										
HEMBA1001581	6.6	2.62	2.74	9.63	8.35	7.34	4.2	4.87	7.29	*	+								
HEMBA1001582	1.39	1.89	0.99	1.46	1.52	1.21	1.87	1.43	1.14										
HEMBA1001585	3.5	1.76	2.06	4.04	4.61	4.34	2.06	2.32	2.78	*	+								
HEMBA1001589	5.07	3.16	2.15	3.41	3.1	3.21	3.05	2.93	3.94										
HEMBA1001595	13.49	4.3	6.8	10.71	10.28	9.89	6.83	8.2	8.67										
HEMBA1001604	5.72	2.28	3.75	6.52	7.03	5.34	2.89	3.22	3.58										
HEMBA1001609	8.03	3.96	3.18	8.15	6.4	9.15	2.3	4.25	3.65										
HEMBA1001615	46.6	32.92	22.49	33.05	34.32	33.44	126.5	104.9	149.7	**	+								
HEMBA1001620	14.48	8.32	7.64	17.62	16.71	15.28	9.29	10.91	11.09	*	+								
HEMBA1001621	9.93	5.63	3.68	7.55	6.93	6.59	5	4.37	6.21										
HEMBA1001635	5.73	3.82	2.42	4.14	4.05	5.67	2.93	3.94	3.69										
HEMBA1001634	4.59	1.44	3.08	3.97	3.02	3.88	3.71	3.49	4.55										
HEMBA1001640	3.49	0.97	1.46	3.57	2.02	3.07	2.4	3.06	2.05										
HEMBA1001647	6.4	2.49	4.47	3.2	5.99	5.63	2.28	4.03	4.16										

8/180

1493

1494

HEMBA1001601	21.79	9.98	12.75	16.31	17.89	15.92	12.62	14.58	15.88						
HEMBA1001605	4.81	3.57	3.37	4.17	7.59	5.82	3.99	4.47	4.7						
HEMBA1001608	2.18	2.11	2.13	1.33	1.53	2.6	1.84	1.15	1.86						
HEMBA1001661	8.45	3.05	2.97	4.66	4.8	5.77	3.88	4.28	4.23						
HEMBA1001665	5.86	2.62	4.27	4.6	3.94	3.51	4.69	4.17	4.52						
HEMBA1001670	4.7	2.38	3.53	6.5	7.04	7.21	4.56	5.94	4.89**	+					
HEMBA1001672	2.9	1.62	1.17	2.74	2.64	2.91	2.23	3.35	2.84						
HEMBA1001673	9.39	2.65	3.57	12.29	9.95	9.16	6.04	3.4	6.06						
HEMBA1001675	2.77	1.09	1.9	3.14	3.42	1.99	2.07	3.04	2.09						
HEMBA1001676	66.2	42	41.28	59.83	62.25	61.28	35.33	41.76	48.98						
HEMBA1001678	23.82	16.82	12.46	26.08	27.44	24.59	15.29	14.2	16.03						
HEMBA1001680	7.67	3.71	3.69	6.51	7.15	6.71	4.41	4.86	5.34						
HEMBA1001681	1.95	0.92	1.52	1.86	1.78	2.38	1.26	2.56	1.49						
HEMBA1001684	10.32	4.07	5.37	13.29	14.64	14.01	8.6	7.77	8.12*	+					
HEMBA1001685	1.84	2.2	0.62	1.62	1.54	2.31	1.72	2.13	0.77						
HEMBA1001702	3.21	1.66	2.35	4.83	3.35	4.17	3.17	4.1	3.6						
HEMBA1001709	3.9	1.96	2.65	5.53	4.06	6.56	5.94	7.83	7.54			**			
HEMBA1001711	2.38	2.81	1.61	5.64	7.85	8.65	3.33	2.8	5.34**	+					
HEMBA1001712	2.87	1.69	2.03	2.84	2.47	3.33	3.23	2.26	2.84						
HEMBA1001714	27.51	15.33	17.22	17.64	16.58	15.17	22.02	17.65	27.85						
HEMBA1001717	1.6	0.57	0.95	1.72	1.13	1.76	8.51	5.96	6.55			**			
HEMBA1001718	3.34	3.04	3.56	7.23	5.88	7.76	3.79	4.78	3.34**	+					
HEMBA1001723	3.28	1.43	2.31	5.16	4.28	5.3	2.9	4.31	2.84*	+					
HEMBA1001731	2.16	1.22	2.13	2.79	1.84	2.37	1.77	2.95	2.23						
HEMBA1001734	2.33	0.57	2.08	3.71	2.97	2.91	2.16	2.87	2.2						
HEMBA1001736	8.5	4.87	4.76	1.77	7.6	9.06	7.56	6.14	10.7						
HEMBA1001741	1.43	1.25	0.91	2.83	2.87	2.84	0.76	1.93	1.43**	+					
HEMBA1001744	1.28	0.91	0.85	1.4	1.01	1.73	0.65	1.88	1.22						
HEMBA1001745	3.12	1.1	1.48	2.46	2.57	2.63	2.55	3.03	2.51						
HEMBA1001746	1.85	2.08	1.47	2.46	2.29	3.39	2.8	3.77	3.54			*			
HEMBA1001761	4.88	2.73	3.04	7.77	5.44	7.35	2.96	5.66	4.32*						
HEMBA1001762	1.84	0.76	1.19	2.52	2.18	2.84	1.18	3.82	1*	+					
HEMBA1001781	3.69	1.25	2.05	4.27	2.77	4.	4.	2.36	3.3	2.22					
HEMBA1001784	5.2	3.84	2.76	3.59	2.92	3.32	3.06	2.91	4.28						
HEMBA1001791	11.2	5.23	3.55	8.42	10	8.96	7.67	7.29	10.19						
HEMBA1001794	16.08	14.18	10.11	17.79	20.03	18.56	11.08	17.68	19.33*	+					
HEMBA1001800	3.13	2.01	2.42	2.99	2.87	3.62	3.26	1.68	2.16						
HEMBA1001803	1.53	0.75	0.44	1.21	1.11	1.38	1.41	1.74	1.3						
HEMBA1001804	13.32	7.17	6.9	11.34	8.08	9.13	6.64	6.79	7.2						
HEMBA1001808	2.99	2.64	1.45	3.65	1.42	1.93	2.32	1.6	1.79						
HEMBA1001809	8.19	6.19	4.29	9.5	5.47	7.96	5.87	6.3	5.81						
HEMBA1001811	22.78	13.64	9.05	9.98	11.16	7.43	11.09	9.04	12.53						
HEMBA1001815	6.31	3.66	3.82	9.75	9.09	7.86	4.84	3.89	4.99*	+					
HEMBA1001816	2.42	2.34	2.86	3.29	2.61	3.73	2.83	1.8	2.99						
HEMBA1001819	6.29	3.74	3.76	9.76	8.91	7.84	4.59	4.2	3.42*	+					
HEMBA1001820	0.7	1.31	0.35	1.28	0.85	0.69	1	1.34	0.88						
HEMBA1001822	14.41	5.14	6.88	15.5	12.29	15.38	7.82	5.44	7.99						
HEMBA1001824	8.95	4.75	6.46	8.72	13.86	15.64	6.51	9.15	7.54						
HEMBA1001835	1.68	1.35	0.6	1.34	1.74	3.48	1.4	1.22	1.48						
HEMBA1001844	7.57	4.41	3.42	10.11	10.97	6.9	6.82	4.22	6.02						
HEMBA1001847	7.9	4.44	3.77	6.2	5.53	6.41	3.82	3.06	3.92						
HEMBA1001849	8.79	2.94	4.27	12.39	11.89	10.26	7.27	5.97	7.41*	+					
HEMBA1001858	7.06	2.55	2.83	6.48	6.52	7.6	4.57	4.3	4.89						
HEMBA1001861	1.79	0.52	1.23	3.92	3.68	4.21	1.63	1.37	1.83**	+					
HEMBA1001862	20.07	14.33	16.51	10.49	11.03	16.77	24.43	27.68	20.89			*			
HEMBA1001864	1.89	1.29	0.85	2.99	3	2.74	1.08	1.87	1.04**	+					

9/180

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【表 179】

1497

1498

HEMBA1002084	1.07	0.5	0.79	1.77	1.77	2.12	1.8	1.72	1.8	**	+	**	+
HEMBA1002085	15.53	10.5	9.09	3.93	5.17	5.54	4.22	4.66	4.34	*	-	*	-
HEMBA1002092	6.36	2.95	3.86	3.82	3.84	2.97	3.77	3.66	5.02				
HEMBA1002098	2.76	1.13	1.61	2.4	2.24	2.53	2.57	2.73	1.55				
HEMBA1002100	32.5	21.44	18.67	25.5	28.16	25.35	18.35	13.17	17.71				
HEMBA1002101	14.23	9.44	8.67	29.88	21.28	21.61	20.63	10.83	13.44	*	+		
HEMBA1002102	5.78	2.45	5.61	10.26	9.25	10.76	5.53	7.91	7.68	**	+		
HEMBA1002105	3.54	2.37	3.22	6.12	5.06	5.65	3.82	6.51	5.09	**	+		
HEMBA1002107	11.45	5.11	6.25	8.68	8.52	8.38	12.57	12.66	17.5				
HEMBA1002113	32.25	19.17	17.4	39.34	45.35	45.81	28.29	21.95	34.31	*	+		
HEMBA1002119	2.11	2.17	0.99	2.79	2.14	2.54	2.06	2.87	1.79				
HEMBA1002125	5.83	2.4	2.92	5.45	9.25	7.16	7.44	6.34	6.72				
HEMBA1002131	5.93	2	3.14	4.14	4.06	4.13	3.5	4.3	3.28				
HEMBA1002133	6.81	5.25	2.52	6.36	5.83	7.36	4.72	7.3	4.48				
HEMBA1002139	1.09	0.26	0.36	1.2	0.84	1.33	0.99	2.43	0.56				
HEMBA1002141	1.29	0.49	1.21	2.38	1.03	1.99	0.5	1.42	1.34				
HEMBA1002144	5.69	3.1	2.06	7.29	6.78	8.63	2.59	3.43	5.33	*	+		
HEMBA1002147	21.38	10.63	10.33	16.26	8.66	14.72	7.7	9.8	14.04				
HEMBA1002150	19.09	10.95	13.29	13.45	10.91	11.19	15.49	16.53	17.44				
HEMBA1002151	5.57	4.52	3.73	5.15	5.43	4.75	6.45	4.35	4.86				
HEMBA1002153	2.06	0.67	0.65	2.43	2.33	1.79	1.41	1.49	1.24				
HEMBA1002156	6.64	2.07	2.79	3.48	2.76	4.92	4.24	4.29	3.26				
HEMBA1002160	9.96	4.66	4.52	11.03	12.78	11.54	5.12	4.86	6.62	*	+		
HEMBA1002161	5.93	2.84	3.76	7.56	5.8	7.54	3.32	4.13	3.25				
HEMBA1002162	7.92	3.54	4.29	9.23	12.27	9.59	6.96	4.68	6.43	*	+		
HEMBA1002163	16.52	8.9	8.29	30.66	23.8	18.1	23.47	24.41	36.58	*	+	*	+
HEMBA1002164	6.58	3.37	3.2	7.61	7.12	6.96	5.68	4.84	5.16				
HEMBA1002166	39.64	27.28	27.86	36.11	45.05	43.8	20.24	20.85	22.71				
HEMBA1002167	4.76	1.86	1.62	2.99	2.78	2.27	3.13	3.05	2.27				
HEMBA1002173	5.99	4.25	4.52	7.86	9.55	7.59	5.43	4.55	6.47	*	+		
HEMBA1002177	7.43	2.78	2.92	3.23	3.61	5.94	3.11	3.88	4.09				
HEMBA1002178	5.72	4.28	4.88	4.38	4.69	4.23	3.54	5.04	4.32				
HEMBA1002179	38.56	31.74	22.53	17.89	19.71	18.71	27.72	23.97	26.16				
HEMBA1002185	6.54	3.16	3.12	9.32	10.15	8.6	6.14	5.78	6.76	*	+		
HEMBA1002188	8.98	4.74	6.37	7.79	6.15	7.58	6.43	5.81	6.6				
HEMBA1002189	3.48	3.26	1.78	4.27	5.47	4.09	2.69	3.88	3.54				
HEMBA1002191	8.3	3.89	4.67	8.84	6.83	6.19	5.91	5.98	6.36				
HEMBA1002192	5.28	4.26	4.29	8.27	6.01	5.9	2.94	2.49	2.82		**	-	
HEMBA1002195	5.98	3.67	4.11	6.21	5.77	4.89	3.93	4.26	3.98				
HEMBA1002196	1.16	1.29	1.53	2.22	2.69	3.34	2.25	2.29	2.94	*	+	**	+
HEMBA1002199	2.9	1.1	2.41	4.59	4.69	3.07	3.88	2.62	3.82				
HEMBA1002204	3.61	1.66	0.98	2.22	2.66	1.99	3.47	1.11	1.87				
HEMBA1002208	48.26	35.92	30.61	48.99	56.44	45.32	18.77	22.83	23.91		*	-	
HEMBA1002212	1.63	2.93	1.64	4.46	4.61	4.63	3.31	1.91	1.67	**	+		
HEMBA1002215	6.24	3.92	3.6	5.45	4.91	5.62	4.3	4.83	3.3				
HEMBA1002217	18.63	10.54	10.96	10.92	19.47	21.75	8.18	9.72	7.73				
HEMBA1002220	2.36	1.42	1.13	2.73	2.21	2.69	1.63	2.43	2.05				
HEMBA1002226	7.06	3.57	4.14	9.44	8.41	9.81	3.48	7.93	5.79	*	+		
HEMBA1002227	23.89	11.28	12.65	63.81	64.96	61.28	43.22	41.38	46.55	**	+	**	+
HEMBA1002229	12.93	9.6	8.96	22.59	17.4	16.24	11.18	9.43	9.71	*	+		
HEMBA1002237	2.73	1.56	1.22	2.88	3.54	2.57	2.65	2.19	1.5				
HEMBA1002239	9.11	4.97	3.45	14.42	9.61	11	4.46	5.84	6.91				
HEMBA1002241	4.16	2.92	3.35	4.29	3.16	4.98	3.45	3.5	3.33				
HEMBA1002253	2	1.21	0.86	1.18	1.75	2.13	1.77	1.87	1.29				
HEMBA1002257	2.5	1.06	1.11	1.47	1.72	1.38	1.46	2.73	1.02				
HEMBA1002259	3.93	2.57	3.46	3.84	3.35	3.79	1.58	3.6	2.24				

11/180

1499

1500

HEMBA1002263	19.33	13.63	11.06	41.08	43.27	39.59	22.08	18	19.52	**	+		
HEMBA1002265	5.77	2.24	2.87	4.81	4.22	4.54	5.24	3.12	3.12				
HEMBA1002267	6.66	4.16	4.1	9.3	9.47	10.3	6.16	5.48	5.21	**	+		
HEMBA1002270	6.34	3.34	3.58	7.78	8.98	7.9	3.76	4.01	4.91	*	+		
HEMBA1002284	2.71	2.63	1.38	2.66	4.03	3.95	2.71	3.54	3.82				
HEMBA1002290	7.29	3.76	4.66	10.41	13.32	10.04	5.97	7.37	8.02	*	+		
HEMBA1002302	11.09	4.74	4.9	14.47	14.16	14.79	5.34	6.14	6.47	*	+		
HEMBA1002304	2.15	1.99	3.2	4.13	2.57	4.4	1.76	2.42	1.31				
HEMBA1002307	20.52	10.07	9.13	9.76	9.15	8.21	13.28	14.24	17.09				
HEMBA1002316	21.96	17.53	15.62	14.66	14.2	13.67	14.54	18.03	17.67				
HEMBA1002319	3.87	2.44	2.95	2.86	3.71	4.51	3	3.92	3.09				
HEMBA1002320	2.67	1.82	1.12	4.11	5.01	6.24	3.84	4.14	3.6	*	+	*	+
HEMBA1002321	1.46	2.38	0.87	3.05	1.97	2.21	1.05	1.18	1.29				
HEMBA1002324	4.66	1.71	1.99	5.92	5.51	4.89	3	3.99	2.69				
HEMBA1002333	4.92	1.14	2.37	2.57	3.45	2.77	2.04	3.2	1.93				
HEMBA1002337	5.38	3.22	4.87	9.22	12.3	11.34	4.19	5.44	4.11	**	+		
HEMBA1002339	23.81	10.43	6.17	11.11	15.1	14.91	11.67	12.27	12.43				
HEMBA1002341	7.39	3.74	4.25	4.55	4.12	3.82	6.09	5.66	5.69				
HEMBA1002348	2.07	1.83	0.9	1.44	1.88	2.08	1.92	2.6	1.34				
HEMBA1002349	1.51	1.42	0.34	1.38	1.3	1.96	1.46	2.19	1.38				
HEMBA1002353	1.79	1.25	2.28	2.64	3.11	3.43	2.11	1.34	1.36	*	+		
HEMBA1002356	13.39	6.02	7.85	8.42	10.26	11	4.88	6.24	6.12				
HEMBA1002357	136.4	89.6	109	142.6	135.4	152.8	57.09	66.8	75.58		*	-	
HEMBA1002360	6.54	3.66	5.93	10.16	10.44	10.51	8.07	9.62	8.15	**	+	*	+
HEMBA1002363	9.05	6.26	4.11	8.4	5.32	7.47	3.78	3.67	4.84				
HEMBA1002365	2.33	1.04	1.69	2.69	1.93	1.79	0.53	1.83	2.11				
HEMBA1002370	2.04	0.84	0.68	5.63	6.49	6.21	1.4	3.02	2.46	**	+		
HEMBA1002374	8.05	4.75	3.85	6.96	7.96	4.55	6.91	5.19	7.37				
HEMBA1002376	22.58	10.7	11.64	20.42	22.01	21.09	9.22	9.95	12.27				
HEMBA1002377	22.23	20.26	24.74	17.13	16.56	16.97	12.65	4.84	13.5	*	*	-	
HEMBA1002380	10.33	4.73	6.12	25.3	20.75	23.1	10.39	11.3	10.43	**	+		
HEMBA1002381	6.11	3.6	4.83	7.07	8.7	10.4	3.87	4.54	4.53	*	+		
HEMBA1002384	15.5	10.84	6.42	29.27	32.78	29.1	8.58	9.53	10.47	**	+		
HEMBA1002389	4.27	1.82	1.04	3.34	2.49	2.48	1.75	2.27	2.21				
HEMBA1002396	5.31	1.45	2.21	3.61	3.86	4.27	4.37	4.75	6.22				
HEMBA1002402	4.83	1.75	1.81	2.54	2.69	3.67	3.46	2.38	3.41				
HEMBA1002417	10.25	4.91	5.09	7.22	6.91	7.47	6.16	5.78	7.28				
HEMBA1002419	5.08	2.09	1.3	5.6	4.81	5.12	3.66	3.43	3.31				
HEMBA1002428	9.17	4.99	7.48	15.98	15.18	16.1	7.35	7.5	9.4	**	+		
HEMBA1002421	3.35	2.15	2.59	6.22	6.03	5.26	6.83	6.92	8.74	**	+	**	+
HEMBA1002423	1.54	0.63	0.83	2.44	3.48	3.88	1.7	2.52	2.63	*	+	*	+
HEMBA1002424	8.4	2.37	3.82	4.43	5.04	5.22	3.68	2.96	3.47				
HEMBA1002426	6.49	3.96	3.42	4.7	3.88	6.41	6.1	4.1	4.43				
HEMBA1002434	2.26	0.52	0.37	1.48	0.95	2.01	1.47	2.19	1.34				
HEMBA1002439	5.88	2.46	3.67	5.4	4.16	4.53	3.95	4.58	3.74				
HEMBA1002441	9.17	5.14	6.07	34.35	24.41	25.8	23.81	16.3	21.56	**	+	**	+
HEMBA1002454	5.79	2.67	3.42	5.87	3.7	3.81	3.97	3.14	3.78				
HEMBA1002459	25.18	17.65	26.81	56.49	54.86	61.69	25.04	31.59	38.07	**	+		
HEMBA1002468	13.9	7.3	5.63	4.27	4.5	4.21	3.8	3.84	3.88				
HEMBA1002462	5.97	3.49	2.63	4.68	5.48	5.1	5.76	6.26	5.62				
HEMBA1002465	1.48	0.35	0.87	1.94	1.91	2	2	1.62	1.54	*	+		
HEMBA1002469	10.61	5.54	6.1	9.43	9.29	9.35	6.49	6.37	7.65				
HEMBA1002475	2.44	1.25	1.2	2.62	1.19	1.93	1.75	2.35	1.4				
HEMBA1002477	4.33	2.21	3.54	6.33	6.45	9.03	4.12	5.33	4.08	*	+		
HEMBA1002486	12.76	7.21	8.4	13.9	9.23	9.97	6.6	9.31	9.9				
HEMBA1002481	4.17	1.44	3.57	5.7	5.97	7.71	3.35	4.98	3.12	*	+		

12/180

1501

1502

HEMBA1002486	8.76	6.38	4.66	8.52	8.8	10.2	5.18	4.82	7										
HEMBA1002489	4.65	2.87	1.43	3.68	2.57	3.08	3.01	1.75	3.98										
HEMBA1002492	3.72	2.75	1.63	4.11	3.81	4.48	2.24	3.36	3.9										
HEMBA1002494	2.75	1.45	1.13	1.68	1.82	1.19	2.23	1.05	1.96										
HEMBA1002498	4.03	2.44	2.73	2.79	3.44	4.73	2.7	2.56	4.15										
HEMBA1002503	5.04	2.61	2.84	6.43	4.88	5.28	3.23	3.79	3.13										
HEMBA1002504	8.07	4.4	4.13	10.71	10.32	10.08	4.47	6.58	5.92	*	+								
HEMBA1002508	5.99	4.98	4.38	8.82	14.4	16.34	4.8	6.77	5.33	*	+								
HEMBA1002513	8.6	4.28	4.52	7.08	4.68	6.71	4.93	3.86	4.51										
HEMBA1002515	4.33	1.73	2.07	3.29	2.16	3.66	2.65	1.63	3.58										
HEMBA1002524	9.35	6	4.75	8.16	6.47	7.51	5.77	5.05	6.67										
HEMBA1002538	4.58	2.03	1.84	2.98	3.05	4.53	2.16	2.92	2.68										
HEMBA1002542	8.07	5.4	5.41	9.41	8.04	9.27	4.65	5.75	5.16										
HEMBA1002544	3.1	1.76	1.69	4.47	3.6	3.68	2.18	2.17	2.61	*	+								
HEMBA1002546	50.52	34.29	29.94	56.51	60.33	61.14	35.34	44.64	38.68	*	+								
HEMBA1002547	2.3	1.72	2.07	1.6	3.25	2.8	2.97	4.34	2.32										
HEMBA1002550	7.14	5.4	3.96	4.54	4.38	4.87	6.51	4.38	5.24										
HEMBA1002551	5.47	2.09	2.27	5.04	4.39	3.41	4.06	3.2	3.87										
HEMBA1002553	12.19	3.86	6.34	10.16	9.24	10.66	6.5	6.73	6.78										
HEMBA1002555	1.98	0.86	1	1.95	2.49	2.76	2.25	1.97	2.82										
HEMBA1002558	7.34	3.99	4.45	10.47	9.14	11.18	5.75	4.9	5.48	*	+								
HEMBA1002561	1.53	2.23	1.45	3.76	4.16	3.85	2.34	2.9	2.42	*	+								
HEMBA1002562	2.58	1.09	1.24	1.55	1.58	1.46	1.13	1.38	1.77										
HEMBA1002568	4.34	2.05	1.84	2.65	3.18	3.63	2.01	3.91	2.77										
HEMBA1002569	10.12	2.96	3.15	6.04	6.91	7.8	6.66	5.49	5.73										
HEMBA1002570	17.18	8.39	8.43	7.74	7.84	6.32	4.15	4.68	4.47										
HEMBA1002574	9.13	5.2	4.08	4.71	4.69	3.46	6.41	4.34	4.04										
HEMBA1002583	2.63	1.94	1.44	4.35	4.76	4.81	4.07	4.23	4.71	**	+	**	+						
HEMBA1002587	9.65	5.73	4.29	5.38	5.09	6.69	6.95	4.55	5.87										
HEMBA1002590	3	2.82	3.17	5.3	7.12	7.9	3.16	4.25	3.45	*	+								
HEMBA1002592	7.22	3.8	5.73	9.2	7.27	11.07	4.7	6.52	5.38										
HEMBA1002595	6.26	2.72	4.83	2.78	4.06	4.2	3.48	5.01	4.73										
HEMBA1002609	4.35	4.09	2.17	4.02	4.01	4.31	3.53	3.64	3.18										
HEMBA1002617	3.95	2.7	1.65	11.81	11.46	11.36	4.49	2.86	3.96	**	+								
HEMBA1002619	6.56	3.72	3.15	6.01	4.48	4.66	4.55	5.76	4.4										
HEMBA1002621	1.33	2.05	0.58	1.87	2.25	1.68	1.25	2.13	1.22										
HEMBA1002624	10.87	5.76	5.5	10.8	7.15	9.93	6.61	9.08	8.33										
HEMBA1002628	2.46	1.89	1.56	8.26	8.9	8.55	6.5	6.76	7.64	**	+	**	+						
HEMBA1002629	2.92	1.59	1.72	2.35	1.78	3.13	2.15	2.22	2.55										
HEMBA1002632	3.01	3.25	2.45	3.55	4.56	5.56	2.28	3.32	2.22										
HEMBA1002645	5.23	3.12	3.3	9.71	9.85	8.47	4.08	4.56	3.84	**	+								
HEMBA1002651	2.74	3.3	3.7	3.63	3.14	3.96	3.35	3.43	3.18										
HEMBA1002652	10.09	4.55	2.8	3.94	7.21	6.43	4.43	4.54	4.59										
HEMBA1002659	10	4.51	4.33	9.97	11.66	8.87	6.05	6.33	5.41										
HEMBA1002661	4.42	2.54	1.79	6.87	7.34	8.27	4.97	3.82	3.85	**	+								
HEMBA1002666	3.37	1.93	1.85	2.66	2.68	3.23	2.59	2.71	2.16										
HEMBA1002667	3.38	2.19	2.12	5.95	5.18	5.02	0.91	3	2.2	**	+								
HEMBA1002673	24.31	16.62	13.86	16.81	24.76	24.88	10.84	15.49	10.36										
HEMBA1002678	6.22	4.52	2.39	8.83	7	9.06	5.27	5.81	5.58	*	+								
HEMBA1002679	6.14	2.98	2.3	7.06	5.91	5.92	4.9	4.03	4.53										
HEMBA1002688	2.43	1.85	1.49	1.28	2.14	1.93	0.91	1.94	1.12										
HEMBA1002696	5.94	3.2	2.68	4.46	5.16	4.1	3.48	3.38	3.28										
HEMBA1002703	14.6	8	9.65	11.88	14.48	12.25	7.74	9.23	10.55										
HEMBA1002706	14.74	6.11	9.63	11.39	13.51	13.79	6.16	6.12	6.47										
HEMBA1002712	5.57	3.22	3.91	7.46	9.02	7.13	2.62	3.73	3.7	*	+								
HEMBA1002715	7.56	4.05	6.71	7.13	9.71	10.17	4.38	6.93	5.48										

13/180

【0654】

【表183】

1503

1504

HEMBA1002716	2.33	1.79	1.1	2.97	1.95	2.33	1.67	1.27	1.06										
HEMBA1002718	16.72	11.81	9.31	17.97	12.96	15.17	10.44	10.19	11.13										
HEMBA1002728	9.67	3.34	5.97	10.6	12.26	15.33	7.76	5.08	9.8	*	+								
HEMBA1002730	7.86	2.52	3.4	5.36	7.91	6.74	6.78	4.96	7.37										
HEMBA1002734	7.73	4.31	3.55	7.93	6.46	7.46	5.62	6.29	6.83										
HEMBA1002741	3.65	1.6	2.01	2.64	2.74	2.6	1.48	2.29	1.92										
HEMBA1002746	6.82	4.06	4.19	4.98	4.66	5.4	2.78	4.2	3.33										
HEMBA1002748	4.16	2.16	3.32	2.33	4.45	4.03	2.42	2.8	3.88										
HEMBA1002750	6.45	3.44	3.09	5.38	6.22	7.28	3.44	2.24	3.97										
HEMBA1002753	6.83	3.3	3.88	9.75	9.18	10.07	4.45	5.29	5.42	*	+								
HEMBA1002759	2.47	0.92	1.55	4.32	3.79	4.12	2.56	2.66	2.65	**	+								
HEMBA1002763	17.79	8.69	9.49	11.93	10.68	12.89	9.46	10.98	11.31										
HEMBA1002767	4.86	3.64	4.15	4.69	4.37	5.27	4.84	4.88	6.63										
HEMBA1002768	7.65	3.89	4.38	7.6	7.36	7.85	6.31	6.75	7.16										
HEMBA1002769	6.55	2.6	4.29	4.3	5.76	5.49	4.08	5.08	4.57										
HEMBA1002770	10.29	6.74	8.19	11.22	11.06	13.4	7.03	6.36	8.42										
HEMBA1002777	9.75	4.7	5.71	8.59	8.79	9.46	6.38	4.3	7.48										
HEMBA1002779	19.22	10.66	6.22	15.16	13.63	10.21	10.37	10.01	10.31										
HEMBA1002780	5.7	2.86	3	6.99	7.8	9.55	4.79	4.73	6.4	*	+								
HEMBA1002780	4.99	2.33	3.07	6.37	8.93	7.96	4.08	3.78	4.9	*	+								
HEMBA1002784	8.37	5.67	4.58	5.78	6.13	8.44	6.79	6.5	6.15										
HEMBA1002798	1.26	0.86	1.65	2.72	2.3	1.86	0.87	2.64	0.77	*	+								
HEMBA1002801	1.99	0.93	1.36	4.21	2.6	1.85	2.71	2.29	3.22										
HEMBA1002810	9.65	4.57	5.68	13.26	12.11	9.75	5.77	6.41	6.28										
HEMBA1002816	9.84	4.52	4.72	9.31	6.58	9.2	5.89	5.54	5.86										
HEMBA1002818	13.95	7.65	7.85	12.57	11.48	11.5	11.94	8.46	10.87										
HEMBA1002820	8.63	4.01	5.8	12.08	16.06	13.75	7.38	6.93	7.73	*	+								
HEMBA1002826	2.06	0.77	0.96	1	0.94	1.69	1.3	2.13	0.88										
HEMBA1002833	9.88	4.57	5.73	7.08	7.89	7.35	7.95	8.57	7.16										
HEMBA1002850	0.76	0.3	1.24	1.8	1.57	1.81	0.67	2.12	1.24	*	+								
HEMBA1002862	2.92	2.24	3.55	9.63	8.86	7.72	5.29	8.86	7.89	**	+	*	+						
HEMBA1002863	3.16	2.79	5.23	4.86	5.55	5.31	3.6	5.86	5.95										
HEMBA1002867	3.74	1.09	1.41	1.95	2.42	2.24	1.51	1.85	1.96										
HEMBA1002876	10.81	3.46	4.85	5.22	5.51	6.47	5.11	4.45	5.37										
HEMBA1002886	1.73	1.14	1.3	1.8	3.11	2.84	1.24	1.52	0.93	*	+								
HEMBA1002896	5.56	2.89	2.26	4.16	5.6	6.36	4.43	4.26	5.28										
HEMBA1002913	6.83	3.41	4.1	6.13	4.56	5.54	4.6	4.46	4.72										
HEMBA1002921	5.09	1.35	3.42	4.01	3.76	3.47	2.82	3.68	1.76										
HEMBA1002924	3.44	1.46	2.03	3.99	2.79	5.07	4.7	2.86	2.66										
HEMBA1002934	19.41	10.56	13.01	28.28	26.9	31.77	13.81	10.62	17.37	**	+								
HEMBA1002935	5.64	2.51	3.1	9.39	9.17	8.78	4.05	4.5	6.44	**	+								
HEMBA1002937	2.94	0.97	1.56	5.32	3.72	3.3	4.25	3.23	5.41			*	+						
HEMBA1002939	5.23	2.26	1.27	6.12	6.22	7.2	3.36	5.43	4.03	*	+								
HEMBA1002944	2.39	1.05	0.97	2.45	2.94	1.89	1.97	1.66	1.79										
HEMBA1002951	4.82	2.48	2.82	6.08	7.02	6.04	4.67	6.63	5.8	*	+								
HEMBA1002954	3.07	1.62	1.21	5.05	3.53	2.74	1.86	3.21	2.77										
HEMBA1002962	4.7	4.71	2.06	11.63	8.54	7.28	2.97	4.52	4.25	*	+								
HEMBA1002968	7.62	3.18	4.17	11.44	8.51	9.98	4.32	4.58	5.82	*	+								
HEMBA1002976	1.55	2.24	2.05	3.8	4.05	2.91	1.8	3.84	2.44	*	+								
HEMBA1002971	2.55	2.17	1.09	2.11	2.8	2.2	1.84	1.44	2.55										
HEMBA1002973	4.7	1.37	2.41	7.46	7.53	5.02	4.19	3.07	3.54	*	+								
HEMBA1002978	4.6	2.2	2.96	5.07	6.26	5.9	2.87	3.98	1.74	*	+								
HEMBA1002981	10.14	3.92	5.05	6.62	5.73	6.85	4.75	3.37	4.22										
HEMBA1002985	5.65	3.15	2.63	4.75	6.26	6.46	4.22	6.1	4.66										
HEMBA1002986	8.06	6.02	4.91	10.22	16.35	15.33	12.26	16.22	12.17	*	+	*	+						
HEMBA1002988	1.58	0.97	1.43	5.23	7.34	7.21	3.78	3.98	3.2	**	+	**	+						

14/180

【0655】

【表184】

1505

1506

HEMBA1002972	9.81	4.26	5.48	8.62	8.82	10.46	6.24	6.81	7.68										
HEMBA1002975	9.95	5.67	5.79	12.67	13.82	15.45	7.42	8.68	5.34	*									
HEMBA1002977	5.35	3.23	2.63	6.04	6.82	4.47	3.67	4.27	4.14										
HEMBA1002979	1.41	1.2		1.77	1.86	2.15	1.32	1.54	1.79	*									
HEMBA1003004	4.4	2.05	2.04	4.44	2.35	3.6	4.34	2.86	3.43										
HEMBA1003006	3.81	3.03	1.95	4.39	5.85	4.42	3.51	4.58	4.16										
HEMBA1003008	3.21	2.19	2.5	3.68	6.17	6.62	2.11	3.8	2.8	*									
HEMBA1003011	7.74	5.2	3.87	9.69	18.49	13.68	7.18	7.68	6.46	*									
HEMBA1003017	2.46	2.25	2.2	3.48	3.21	5.26	3.71	4.99	5.27			**	+						
HEMBA1003019	16.49	15.58	12.66	14.01	22.6	13.51	9.84	22.76	21.22										
HEMBA1003031	7	6.8	4.83	11.72	14.51	12.51	5.21	5.97	6.1	**	+								
HEMBA1003033	8.54	5.52	5.51	6.83	9.05	7.67	7.01	6.8	7.13										
HEMBA1003033	13.69	8.92	7.92	18.19	20.22	19.59	7.08	10.97	9.51	**	+								
HEMBA1003034	10.16	6.76	5.59	16.34	16.21	18.88	7.61	9.38	7.94	**	+								
HEMBA1003035	0.86	0.59	0.52	1.61	1.97	0.55	0.09	2.49	0.47										
HEMBA1003037	14.14	5.43	5.96	7.58	8.71	8.97	7.73	6.56	7.19										
HEMBA1003041	13.54	5.42	7.39	11.23	11.7	11.68	7.62	7.38	7.89										
HEMBA1003046	10.88	8	7.65	10.34	12.65	9.57	6.83	7.13	6.72										
HEMBA1003047	6.06	2.52	2.2	4.15	3.03	5.74	4.14	4.89	5.07										
HEMBA1003048	4.06	2.13	2.64	5.2	6.24	5.07	5.54	7.31	7.12	*	+	**	+						
HEMBA1003044	1.85	0.88	1.11	2.44	3.01	2.83	0.75	2.53	1.61	*	+								
HEMBA1003047	3.99	3.73	3.24	6.04	4.55	5.67	2.42	3.41	2.73	*	+								
HEMBA1003071	4.89	2	2.46	3.09	3.36	3.95	2.75	4.15	2.46										
HEMBA1003072	5	3.54	3.49	9.31	7.84	7.21	4.62	3.3	3.28	**	+								
HEMBA1003076	17.78	7.65	8.23	14.31	15.74	15.39	9.87	12.41	9.92										
HEMBA1003077	2.58	1.45	1.89	1.93	2.25	1.91	1.66	2.15	1.57										
HEMBA1003078	2.54	1.51	1.55	3	4.23	3.44	2.66	2.38	2.18	*	+								
HEMBA1003079	1.91	1.85	1.65	2.48	2.95	4.35	2.42	3.49	4.71										
HEMBA1003083	3.9	3.64	3.64	4.53	10.29	6.35	3.33	5.25	3.51										
HEMBA1003086	4.22	1.35	2.59	5.79	6.56	7.76	2.96	3.81	3.22	*	+								
HEMBA1003090	4.24	1.39	3.28	3.62	3.38	4	2.6	2.81	2.75										
HEMBA1003094	7.91	4.48	3.84	6.39	5.65	6.37	3.63	7.81	6.29										
HEMBA1003096	2.55	1.26	1.42	4.86	4.02	4.86	6.22	5.89	7.19	**	+	**	+						
HEMBA1003098	13.1	7.22	6.89	14.21	8.08	14.42	7.57	4.87	3.41										
HEMBA1003101	3.86	1.85	2.21	3.42	3.01	2.84	5.05	3.05	3.57										
HEMBA1003109	4.5	2.81	2.78	4.25	4.4	4.14	3.32	3.04	3.75										
HEMBA1003114	4.72	1.49	2.76	4.85	5.37	3.83	2.72	3.02	2.47										
HEMBA1003117	3.34	1.32	1.84	1.94	3.48	3.4	1.15	2.8	1.47										
HEMBA1003120	6.26	3.04	4.46	8.53	11.03	9.87	2.19	4.52	4.23	*	+								
HEMBA1003129	2.92	2.45	1.66	4.57	4.93	4.76	2.42	2.95	3.22	**	+								
HEMBA1003133	3.76	2.75	2.66	4.15	4.94	3.64	3.03	3.83	3.17										
HEMBA1003136	10.1	5.38	5.56	4.76	5.69	3.55	4.84	6.06	5.16										
HEMBA1003142	3.63	2.31	2.57	5.7	6.12	5.75	4.06	4.52	4.11	**	+	*	+						
HEMBA1003148	3.76	1.84	1.85	6.57	6.35	7.13	3.78	3.2	3.41	**	+								
HEMBA1003151	3.06	1.21	2.06	3.57	3.12	3.47	1.14	2.71	1.88										
HEMBA1003153	0.94	1.17	1.24	1.37	1.78	3.11	1.18	1.39	0.96										
HEMBA1003157	5.21	1.69	3.38	3.87	2.5	3.42	0.86	1.61	0.57										
HEMBA1003166	16.26	12.43	11.19	32.39	35.71	31.69	16.79	24.31	19.76	**	+								
HEMBA1003171	2.89	0.72	0.57	1.31	1.88	0.92	1.38	1.84	1.16										
HEMBA1003175	2.6	1.51	1.44	3.64	3.88	4.37	1.98	2.4	2.06	**	+								
HEMBA1003179	4.43	2.72	4.24	3.15	2.78	4.29	3.01	4.28	3.14										
HEMBA1003186	8.23	6.45	5.52	11.84	15.62	13.23	6.12	9.01	7.56	**	+								
HEMBA1003196	5.41	2.8	3.15	4.6	6.06	5.01	3.65	4.6	3.58										
HEMBA1003197	1.16	0.72	0.59	1.88	2.27	1.84	1.37	1.38	0.69	**	+								
HEMBA1003199	2.2	0.82	0.97	3.9	3.83	3.59	1.11	2.24	1.33	**	+								
HEMBA1003202	6.51	4.3	4.72	9.3	10.18	10.03	5.76	4.86	4.67	**	+								

15/180

【0656】

【表185】

1507

1508

HEMBA1003204	4.47	2.88	1.95	6.42	9.31	7.19	4.16	4.11	3.52	*	+		
HEMBA1003210	5.3	3.32	2.57	8.14	10.48	8.37	17.93	9.12	16.05	**	+	*	+
HEMBA1003212	10.06	5.15	6.84	12.91	14.8	18.49	9.01	8.88	9.76	*	+		
HEMBA1003218	1.85	0.63	1.04	1.36	1.25	1.72	1.4	2.5	1.89				
HEMBA1003220	27.66	24	25.44	26.62	36.09	37.79	16.07	14.85	17.5		**	-	
HEMBA1003222	2.88	1.72	3.36	3.75	3.58	3.59	2.57	3.59	2.87				
HEMBA1003225	2.92	1.48	2.59	3.07	2.81	2.57	2.42	3.81	3.48				
HEMBA1003229	3.63	1	0.92	4.49	4.02	4.36	4.86	6.18	8.35		*	+	
HEMBA1003230	4.81	1.33	1.59	3.63	3.48	2.96	4.65	4.11	4.45				
HEMBA1003235	4.25	2.83	2.72	4.77	5.98	6.44	3.15	3.65	2.94	*	+		
HEMBA1003236	2.61	2.12	2.62	4.85	3.24	5.32	5.66	4.6	3.8	*	+	*	+
HEMBA1003250	1.73	0.34	1.4	2.93	3	2.03	1.83	2.23	1.38	*	+		
HEMBA1003252	5.88	2.96	5.36	7.78	7.79	7.89	4.58	5.63	5.99	*	+		
HEMBA1003257	4.93	1.49	3.88	3.03	4.82	4.08	2.99	3.59	4.04				
HEMBA1003268	0.75	0.26	0.6	2.39	1.18	1.2	0.42	1.31	0.41				
HEMBA1003273	3.46	2.51	1.67	5.94	6.01	5.04	2.19	3.45	3.47	**	+		
HEMBA1003276	1.81	1.29	0.96	4.38	4.69	4.83	2.14	2.73	3.03	**	+	*	+
HEMBA1003277	2.81	1.68	0.99	2.59	2.91	2.66	2.69	2.74	1.67				
HEMBA1003278	1.65	0.9	1.98	2.98	3.92	3.95	2.37	3.01	2.17	**	+		
HEMBA1003280	3.32	1.78	3	4.76	3.3	3.57	2.93	5.18	3.65				
HEMBA1003281	4.06	0.91	2.42	3.46	3.32	3.57	2.53	4.81	3.88				
HEMBA1003284	0.48	0.51	0.58	2.22	0.82	1.41	1.13	2.8	1.15				
HEMBA1003286	3.88	2.4	2.73	5.92	5.88	3.67	2.08	4.79	3.77	*	+		
HEMBA1003291	2.38	1.74	0.96	2.57	3.95	3.8	2.72	4.5	5.97	*	+		
HEMBA1003294	5.2	3.14	3.02	8.15	7.24	7.54	4.43	4.64	6.12	**	+		
HEMBA1003296	3.52	1.49	1.47	1.62	2.44	1.83	2.01	2.49	2.5				
HEMBA1003304	1.33	0.87	0.46	1.14	1.8	1.15	0.92	1.05	1.59				
HEMBA1003306	4.82	1.91	2.68	6.16	5.24	6.21	5.8	5.67	6.01	*	+	*	+
HEMBA1003309	0.64	0.18	0.96	3.28	3.28	2.43	1.17	2.04	1.94	**	+	*	+
HEMBA1003314	30.47	18.15	16.33	19.29	25.08	19.75	20.31	20.79	24.11				
HEMBA1003315	10.03	5	5.86	8.82	6.71	8.85	7.02	6.3	8.18				
HEMBA1003322	6.46	2.81	4.38	11.92	11.23	7.71	5.2	4.77	6.83	*	+		
HEMBA1003326	4.18	1.78	2.35	2.75	2.55	2.84	2.28	3.1	3.12				
HEMBA1003327	1.82	3.14	1.29	2.95	3.45	3.27	2.29	2.03	3.08				
HEMBA1003328	4.01	4	2.1	5.29	8.09	6.1	3.75	5.53	3.53	*	+		
HEMBA1003330	11.21	6.43	6.46	11.55	10.31	11.11	5.39	5.56	6.86				
HEMBA1003348	5.75	4.37	3.56	10.47	9.44	9.51	4.42	5.42	4.82	**	+		
HEMBA1003349	3.52	2.39	2.06	5.95	6.68	6.94	3.15	4.91	3.36	**	+		
HEMBA1003370	20.51	11.56	11.02	25.15	23.1	21.13	12.45	14.72	17.99				
HEMBA1003373	3.04	1.4	0.86	3.17	2.01	3.32	2.12	1.4	2.16				
HEMBA1003376	11.18	5.54	7.92	20.96	23.88	21.25	10.64	10.28	11.12	**	+		
HEMBA1003380	2.3	1.46	1.33	2.34	1.63	1.87	2.49	1.54	2.46				
HEMBA1003384	2.29	0.73	1.56	3.93	3.22	3.27	1.72	2.42	3.12	*	+		
HEMBA1003387	1.34	0.55	1.92	1.88	0.47	0.99	1.2	0.99	1.1				
HEMBA1003392	8.27	4.38	5.55	5.24	7.99	8.63	5.42	7.97	6.53				
HEMBA1003395	1.96	1.22	1.19	2.43	3.54	3.02	1.99	5.5	1.02	*	+		
HEMBA1003399	3.58	3.74	3.33	5.08	4.37	5.04	3.4	3.1	3.67				
HEMBA1003408	10.74	5.28	6.5	8.13	8.07	5.69	7.43	7.79	7.28				
HEMBA1003402	4.66	2.07	1.57	4.25	3.02	2.77	2.27	1.71	2.18				
HEMBA1003403	4.57	4.91	4.99	4.14	4.8	3.26	2.96	3.55	2.8	**	-		
HEMBA1003408	10.68	7.13	5.44	7.16	7.17	7.67	7.62	9.08	7.52				
HEMBA1003412	6.57	4.94	4.07	6.42	6.27	6.69	3.99	6.63	4.67				
HEMBA1003417	4.27	2.26	3.09	1.9	2.03	2.19	2.24	2.99	2.76				
HEMBA1003418	10.03	4.9	6.22	10.24	12.15	12.3	3.46	6.64	4.53				
HEMBA1003420	1.52	0.53	0.73	7.04	2.2	5.31	11.33	12.68	10.88	**	+		
HEMBA1003425	1.37	1.11	1.09	2.68	2.01	2.17	1.69	0.88	0.88	**	+		

16/180

【0657】

【表186】

1509

1510

[illegible]

1511

1512

HEMBA1003466	1.38	1.05	0.83	1.72	1.7	1.06	0.87	1.13	0.89										
HEMBA1003467	14.71	11.01	9.94	14.75	22.82	18.78	15.24	10.59	14.49										
HEMBA1003470	0.91	0.22	0.29	1.11	1.61	1.82	0.56	1.43	0.85	*	+								
HEMBA1003474	26.03	18.94	18.61	21.47	28.7	30.08	14.76	19.25	20.49										
HEMBA1003477	3.73	1.52	2.36	7.63	8.16	6.96	10.74	10.88	8.28	**	+	**	+						
HEMBA1003479	1.48	0.67	1.25	5.41	5.58	4.44	1.61	3.27	2.24	**	+								
HEMBA1003480	6.18	3.86	3.32	4.89	3.65	4.22	2.45	3.41	4.34										
HEMBA1003484	3.07	3.42	2.52	4.93	3.87	3.53	2.61	2.37	4.26										
HEMBA1003490	8.67	4.5	4.89	6.53	3.61	6.33	6.11	7.01	7.57										
HEMBA1003492	6.51	4.39	2.76	7.65	13.21	11.37	6.71	7.24	6.09	*	+								
HEMBA1003782	7.49	3.3	2.54	5.23	6.69	4.84	4.77	3.72	5.73										
HEMBA1003711	5.86	2.58	3.21	3.28	5.31	5.99	2.91	4.08	4.68										
HEMBA1003714	4.3	2.42	1.47	3.54	3.98	3.51	1.5	2.8	3.08										
HEMBA1003715	5.16	2.24	2.94	8.09	8.13	8	2.66	4.48	4.1	**	+								
HEMBA1003717	3.17	2.29	1.96	4.19	3.55	5.52	1.88	1.44	1.67	*	+								
HEMBA1003720	1.56	1.73	1.27	3.11	3.53	3.49	2.3	1.66	3.08	**	+								
HEMBA1003725	1.46	0.94	0.92	3.84	2.37	2.61	2.1	1.7	2.25	*	+	*	+						
HEMBA1003728	6.2	3.24	4.06	5.16	6.27	6.67	5.85	4.48	3.55										
HEMBA1003729	3.99	1.42	2.32	6.36	5.84	4.38	3.64	4.72	3.3	*	+								
HEMBA1003733	1.63	1.1	1	3.52	2.12	1.25	0.95	1.54	1.47										
HEMBA1003733	2.5	4.71	1.16	4.86	6.33	5.47	2.90	3.73	3.5										
HEMBA1003742	6.12	2.9	4.2	5.24	4.87	5.32	2.62	6.27	5.03										
HEMBA1003743	2.64	1.63	1.2	2.32	2.37	3.69	2.34	1.46	1.92										
HEMBA1003758	5.8	2.98	4.74	10.06	11.45	11.44	7.34	3.52	6.11	**	+								
HEMBA1003760	5.32	2.29	2.62	4.55	3.7	4.58	3.57	4.5	4.37										
HEMBA1003764	5.57	1.67	3.47	5.12	2.71	2.62	3.98	3.91	4.87										
HEMBA1003769	11.09	7.81	6.22	7.38	7.99	10.09	8.32	6.25	8.19										
HEMBA1003773	4.06	2.15	2.74	3.4	2.78	2.89	3.34	3.66	3.74										
HEMBA1003783	5.9	5.63	5.15	7.21	10.97	7.92	4.02	5.97	4.35										
HEMBA1003784	1.56	0.55	0.26	1.01	1.64	1.14	0.84	1.59	1										
HEMBA1003794	22.02	14.74	15.29	16.32	23.57	18.51	19.15	20.16	23.83										
HEMBA1003799	3.18	0.83	0.69	1.6	1.44	2.62	1.76	1.29	1.44										
HEMBA1003803	5.18	3.99	2.9	7.41	7.07	8.96	7.06	6.07	5.89	*	+	*	+						
HEMBA1003804	4.31	3.24	3.27	5.11	3.19	3.64	3.81	3.96	3.99										
HEMBA1003805	9.07	8.11	9.22	15.23	14.63	10.98	7.34	10.04	6.52	*	+								
HEMBA1003807	2.26	0.57	1.05	1.41	1.99	1.42	1.14	1.69	1.19										
HEMBA1003810	2.67	2.32	0.99	3.03	2.59	2.69	2.81	4.61	3.57										
HEMBA1003827	25.92	18.96	19.46	14.46	20.55	25.66	14.02	29.91	19.07										
HEMBA1003836	9.8	5.94	7.41	16.46	20.73	18.84	10.1	7.05	9.1	**	+								
HEMBA1003838	29.21	22.41	20.25	35.45	47.13	35.6	26.78	26.74	23.31	*	+								
HEMBA1003843	8.31	5.73	4.45	4.63	2.15	3.75	3.72	2.7	3.17										
HEMBA1003846	26.28	20.72	18.37	21.86	22.27	12.11	9.99	15.1	13.9			*	-						
HEMBA1003854	3.23	2.48	1.56	1.62	2.7	2.03	1.6	2.63	2.09										
HEMBA1003857	5.6	3.94	4.15	8.14	11.16	11.16	4.4	7.01	4.61	**	+								
HEMBA1003864	4.85	1.81	2.77	3.23	4.07	4.12	3.31	3.39	2.74										
HEMBA1003866	1.42	0.62	1.37	1.22	1.21	1.69	0.7	1.76	0.89										
HEMBA1003868	13.28	7.75	6.42	9.42	7.15	9.18	5.91	7.86	7.2										
HEMBA1003879	2.14	1.42	1.52	4.08	4.35	3.36	3.08	2.74	2.28	**	+	*	+						
HEMBA1003889	4.68	2.16	2.83	4.05	4.87	3.64	3.32	3.39	3.6										
HEMBA1003884	5.74	3.71	3.92	4	4.49	5.17	4.44	4.54	4.54										
HEMBA1003883	10.32	6.22	7.2	14.27	16.75	16.47	8.29	8.14	8.72	**	+								
HEMBA1003887	5.7	2.76	3.69	4.8	4.75	6.26	4.28	5.31	3.98										
HEMBA1003898	5.76	2.14	5.85	3.21	4.35	4.1	2.43	1.67	2.34										
HEMBA1003893	24.48	11.91	15.58	30.06	36.65	30.84	15.38	16.42	17.2	*	+								
HEMBA1003896	19.51	13.33	11.04	17.41	24	18.91	15.03	15.04	20.6										
HEMBA1003902	8.4	6.56	5.19	9.58	9.98	7.89	5.71	4.62	5.77										

18/180

1513

1514

HEMBA1003904	2.78	1.45	1.43	2.77	2.17	1.59	1.47	2.66	1.69										
HEMBA1003908	1.69	1.16	1.22	2.42	2.58	2.06	1.92	2.64	1.46	*	+								
HEMBA1003926	72.36	45.24	46.72	61.73	49.96	64.94	25.26	18.43	24.45		*	-							
HEMBA1003937	3.1	1.85	1.98	6.12	8.5	7.61	2.66	5.69	3.16	**	+								
HEMBA1003939	1.28	1.62	1.87	1.85	4.47	4.22	0.72	2.97	2.45										
HEMBA1003940	2.82	0.88	1.71	2.17	3.19	2.37	0.51	2.52	1.7										
HEMBA1003941	4.35	2.77	1.79	1.96	4.65	3.03	2.55	3.88	2.82										
HEMBA1003942	2.44	1.82	1.09	3.64	3.65	2.58	2.38	2.63	2.03	*	+								
HEMBA1003945	9.46	3.83	5.74	8.44	8.96	9.42	7.88	6.57	7.46										
HEMBA1003949	2.14	1.99	0.59	2.89	3.58	3.78	1.92	2.25	1.36	*	+								
HEMBA1003950	1.45	1.52	0.64	1.83	1.87	1.76	1.11	1.8	1.56										
HEMBA1003953	1.96	0.44	1.34	3.08	3.28	3.34	1.95	3.21	1.37	*	+								
HEMBA1003958	6.58	4.78	4.74	10.87	13.86	10.68	4.23	6.52	6.23	**	+								
HEMBA1003959	2.84	3.02	3.46	6.74	9.97	6.27	2.64	3.74	2.94	*	+								
HEMBA1003960	7.33	2.27	2.98	3.59	5.1	3.92	2.8	3.92	3.79										
HEMBA1003966	4.91	3.07	2.16	3.5	4.6	3.28	2.1	2.93	3.48										
HEMBA1003967	5.85	3.63	2.68	3.94	3.8	3.19	1.89	3.17	2.35										
HEMBA1003968	3.76	2.02	2.13	4.21	6.16	3.59	4.13	4.11	3.84										
HEMBA1003974	41.47	29.67	25.73	95.3	104.1	103.5	100	82.53	110.2	**	+	**	+						
HEMBA1003976	2.48	1.1	1.38	2.13	2.25	2.22	1.34	1.82	1.6										
HEMBA1003977	2.19	1.38	1.4	2.42	3.02	1.57	1.86	1.96	1.93										
HEMBA1003978	2.44	1.5	1.92	3.24	3.34	3.83	1.9	2.67	2.77	*	+								
HEMBA1003981	7.98	4.15	3.07	6.67	7.3	7.05	6.37	6.68	8.81										
HEMBA1003982	6.94	4.75	3.19	18.33	22.13	23.04	19.29	21.74	19.78	**	+	**	+						
HEMBA1003983	2.27	1.26	0.95	3.01	1.91	1.85	1.02	2.35	1.03										
HEMBA1003987	3.79	1.42	2.2	4.67	5.44	5.59	3.67	4.19	3.44	*	+								
HEMBA1003989	2.32	1.65	1.59	4.16	4.13	5.73	3.24	3.75	3.69	**	+	**	+						
HEMBA1004000	1.83	1.8	1.37	4.32	4.14	3.96	2.63	3.55	2.34	**	+	*	+						
HEMBA1004006	1.37	0.24	1.22	2.13	0.94	1.32	1.17	2.2	0.95										
HEMBA1004007	6.04	2.39	4.27	10.96	12.86	11.45	4.4	6.87	6.82	**	+								
HEMBA1004010	2.94	1.19	1.2	2.7	3.56	3.85	6.4	6.08	6.71			**	+						
HEMBA1004011	1.7	0.78	0.96	2.15	2.36	1.93	1.31	2.48	1.53	*	+								
HEMBA1004012	3.28	1.3	2.31	6.01	4.99	6.51	2.69	3.48	2.84	**	+								
HEMBA1004013	2.75	2.05	2.56	5.11	5.22	4.78	3.43	3.89	3.43	**	+	*	+						
HEMBA1004024	5.55	4.27	3.76	12.33	16.73	14.13	7.3	6.98	6.78	**	+	*	+						
HEMBA1004029	4.41	3.27	3.73	8.06	10.91	6.74	3.3	6.61	3.72	*	+								
HEMBA1004038	2.95	1.3	1.87	2.87	2.27	1.5	1.15	4.52	1.31										
HEMBA1004042	0.26	0.07	0.48	1.39	1.07	0.6	0.74	2.16	0.79										
HEMBA1004043	1.3	0.85	0.68	0.88	3.2	1.57	1.01	1.7	0.88										
HEMBA1004049	7.55	3.12	4.4	8.61	11.26	7.66	7.55	7.12	7.08										
HEMBA1004049	1.17	0.64	0.7	2.26	3.05	2.36	1.86	2.62	1.77	**	+	*	+						
HEMBA1004051	4.38	1.98	1.73	3.51	4.36	4.02	9.79	8.74	8.25			**	+						
HEMBA1004053	8.63	5.44	4.46	17.89	9.59	13.59	14.13	13.15	12.59			**	+						
HEMBA1004055	2.65	0.36	1.81	2.57	2.89	2.7	1.3	3	1.6										
HEMBA1004056	7.5	3.93	5.65	20.02	18.97	17.23	8.27	9.56	8.48	**	+	*	+						
HEMBA1004060	0.07	0.43	1.07	1.82	1.29	1.56	0.47	1.74	1.31	*	+								
HEMBA1004061	14.25	5.05	4.22	4.44	4.07	3.17	2.96	3.91	3.1										
HEMBA1004067	9.19	5.05	5.06	7.71	7.13	7.12	4.94	6.2	7.71										
HEMBA1004071	14.49	9.52	7.51	14.12	12.05	13.02	8.33	9.62	10.66										
HEMBA1004074	7.06	2.77	2.38	5.08	5.28	4.16	4.21	4.55	5.63										
HEMBA1004078	11.34	10.72	8	8.83	11.03	9.34	8.26	9.99	8.73										
HEMBA1004083	3.75	2.6	2.94	3.51	4.46	4.01	3.83	4.11	2.49										
HEMBA1004084	9.29	6.04	5.26	10.09	10	10.42	3.43	2.6	4.21										
HEMBA1004097	2.9	2.64	1.83	1.78	5.31	3.52	1.65	4.02	2.63										
HEMBA1004100	5.05	2.67	3.16	5.99	4.86	5.28	5.44	5.79	5.01										
HEMBA1004103	10.13	4.33	3.51	10.84	11.41	10.36	6.76	6.57	8.64										

19/180

【0660】

【表189】

1517

1518

HEMBA1004329	6.64	4.05	3.69	10.1	11.36	10.31	6.59	6.39	7.21**	+		
HEMBA1004330	3.08	1.92	1.4	2.71	3.14	3.34	2.44	3.02	3.26			
HEMBA1004334	3.9	1.95	1.91	3.91	3.51	4.57	2.69	2.68	1.57			
HEMBA1004335	4.91	2.24	2.81	7.41	9.8	7.49	5.19	6.78	5.7*	+		
HEMBA1004341	6.84	4.77	5.5	4.53	6.2	3.05	5.02	6.83	6.93			
HEMBA1004344	17.75	13.13	14.74	15.95	19.67	19.67	18.23	19.51	24.66			
HEMBA1004347	4.63	3.35	2.01	5.16	6.48	5.36	2.73	3.19	3.71			
HEMBA1004349	8.89	2.46	3.99	12.23	16.69	10.37	7.71	6.98	8.61*	+		
HEMBA1004353	5.41	3.1	3.3	7.91	8.12	10.45	4.93	5.52	5.36*	+		
HEMBA1004353	8.35	7.6	6.31	15.51	16.38	15.96	6.75	8.35	9.31**	+		
HEMBA1004354	4.38	1.54	2.32	5.25	5.81	6.37	3.27	4.92	3.61*	+		
HEMBA1004356	2.81	2.85	3.03	5.06	4.66	5.46	5.28	5.77	4.17**	+	+	
HEMBA1004360	5.79	2.16	5.01	6.93	5.95	3.72	3.15	6.55	5.08			
HEMBA1004366	2.78	2.3	2.86	5.4	6.73	4.61	2.18	3.01	3.38*	+		
HEMBA1004372	0.38	0.77	0.43	0.47	0.63	0.99	0.52	0.83	0.34			
HEMBA1004377	7.38	3.14	3.85	11.65	12.1	15.48	9.22	8.78	11.95**	+	+	
HEMBA1004389	18.67	11.71	10.38	8.69	8.39	17.15	9.23	8.15	7.38			
HEMBA1004391	2.93	2.48	2.45	7.42	5.09	7.12	3.62	4.64	3.41**	+	+	
HEMBA1004393	18.44	14.15	13.12	19.38	17.77	18.16	22.31	14.59	20.28			
HEMBA1004394	1.18	1.11	1.72	2.3	1.6	2.38	1.09	4.42	1.46			
HEMBA1004396	1.79	1.02	1.22	3.41	3.48	3.73	1.3	3.02	1.73**	+		
HEMBA1004401	4.73	3.38	4.96	4.16	4.54	5.13	2.63	5.44	3.27			
HEMBA1004405	3.95	2.13	1.81	6.15	8.26	6.59	3.78	4.33	5.63**	+		
HEMBA1004408	5.72	3.65	3.17	5.44	6.45	4.46	2.34	3.68	2.97			
HEMBA1004414	8.38	4.86	5.28	9.94	19.52	21.58	6.98	7.48	7.61*	+		
HEMBA1004429	3.38	2.07	1.78	8.58	8.61	9.23	4.27	3.18	4.51**	+		
HEMBA1004433	1.82	1.56	1.04	5.34	5.56	5.46	1.92	2.85	2.38**	+		
HEMBA1004440	2.19	0.58	1.67	2.76	2.16	2.15	1.08	2.89	1.62			
HEMBA1004444	4.28	2	2.33	6.71	7.29	10.11	5.5	5.93	3.39*	+		
HEMBA1004446	1.19	0.41	1.18	2.01	2.51	2.6	0.58	1.63	1.83*	+		
HEMBA1004451	4.92	5.14	2.78	5.62	4.16	5.1	2.95	3.75	4.07			
HEMBA1004452	1.45	1.3	0.96	7.34	8.28	11.36	3.26	5.07	5.69**	+	+	
HEMBA1004454	2.73	3.17	2.58	3.68	3.73	5.7	3.62	3.63	3.66			
HEMBA1004460	8.77	5.29	4.63	9.49	11.6	11.51	5.17	5.78	6.91*	+		
HEMBA1004461	3.02	1.29	1.56	1.22	2.06	2.62	1.48	2	2.51			
HEMBA1004468	9.69	5.12	5.83	5.76	9.08	12.25	6.18	7.22	5.91			
HEMBA1004479	5.17	2.6	2.53	3.06	4.8	5.24	1.98	4.08	3.44			
HEMBA1004482	2.81	3.98	3.7	2.47	3.92	2.52	2.59	2.29	3.11			
HEMBA1004491	1.37	1	0.98	1.25	1.97	1.96	0.89	1.47	2.84			
HEMBA1004499	6.22	5.75	3.57	9.95	9.17	8.62	6.22	6.62	6.45*	+		
HEMBA1004502	3.1	2.59	1.77	4.11	5.34	4.51	4.17	2.98	4.03*	+		
HEMBA1004505	4.8	2.59	1.93	2.42	4.25	3.38	2.91	2.43	2.1			
HEMBA1004506	2.39	1.28	1.21	2.96	3.46	3.27	2.23	2.51	1.92*	+		
HEMBA1004507	70.44	39.05	46.26	43.39	51.75	50.62	19.17	24.55	22.93			
HEMBA1004509	5.46	3.62	4.71	3.53	4.82	5.37	2.96	3.83	2.3			
HEMBA1004513	1.41	0.73	0.59	1.16	1.53	1.37	1.32	1.24	1.24			
HEMBA1004528	3.19	1.97	1.1	3.38	4.01	3.33	4.31	3.09	4.88			
HEMBA1004534	6.12	2.73	4	6.77	8.18	7.93	6.04	5.56	6.21*	+		
HEMBA1004536	4.76	3.38	3.05	3.55	4.6	4.52	2.5	2.23	2.99			
HEMBA1004538	21.21	15.5	13.77	31.9	33.44	32.76	19.4	20.15	17.02**	+		
HEMBA1004542	2.99	2.19	1.59	3.03	3.58	3.02	3.51	3.43	2.25			
HEMBA1004552	7.56	6.12	5.53	7.59	13.46	14.87	4.88	6.44	7.28			
HEMBA1004554	2.07	2.28	0.95	2.8	2.16	2.43	2.95	3.09	3.26			
HEMBA1004558	11.57	6.62	6.21	7.21	8.48	8.56	6.35	6.8	7.65			
HEMBA1004560	4.78	3.27	2.78	3.55	5.31	4.2	5.01	4.88	3.58			
HEMBA1004564	7.43	4.79	5.05	12.74	14.02	11.12	5.94	7.39	6.24**	+		

21/180

1519

1520

HEMBA1004566	28.53	23.96	23.72	13.72	19.42	18.07	12.15	16.7	14.58	*	-	**	-
HEMBA1004573	2.19	1.72	1.51	3.93	5.22	5.71	3.32	3.47	1.99	**	+		
HEMBA1004576	2.94	1.45	1.92	18.03	33.01	34.57	7.81	8.41	10.12	**	+	**	+
HEMBA1004577	5	2.83	2.54	7.07	10.98	8.6	4.13	8.82	4.99	*	+		
HEMBA1004586	5.72	3.41	4.19	8.7	7.46	11.19	4.1	6.11	4.48	*	+		
HEMBA1004596	4.81	2.28	2.02	2.96	3.67	3.46	2.47	2.61	3				
HEMBA1004604	6.48	4.01	3.96	4.74	6.55	5.9	8.49	6.15	4.83				
HEMBA1004607	3.7	2.23	1.35	4.64	5.86	4.48	2.81	3.79	3.93	*	+		
HEMBA1004610	4.03	2.57	2.33	4.52	5.94	4.83	3.02	2.81	3	*	+		
HEMBA1004617	2.21	4.92	1	2.84	9.03	3.69	1.85	2.86	2.94				
HEMBA1004622	5.45	3.28	2.52	5.48	8.39	9.1	4.14	4.48	4.68				
HEMBA1004626	4.11	2.56	2.25	5.1	4.71	5.91	2.73	4.36	3.32	*	+		
HEMBA1004629	3.07	1.77	1.42	3.68	3.77	4.82	1.19	3.75	1.18	*	+		
HEMBA1004631	1.43	2.39	0.95	2.12	1.94	2.84	2.88	1.6	2.44				
HEMBA1004632	2.27	1.83	1.79	2.78	2.92	1.76	2.34	3.5	2				
HEMBA1004633	7.83	5.66	4.81	4.47	6.1	5.15	5.55	4.15	5.55				
HEMBA1004636	6.11	4.03	3.37	5.56	5.52	5.5	4.94	4.1	4.16				
HEMBA1004637	3.8	2.43	1.85	2.17	3.96	3.28	2.95	2.5	2				
HEMBA1004638	1.58	0.7	0.19	0.85	2.26	3.04	1.06	1.19	1.64				
HEMBA1004645	4.58	1.72	2.46	3.58	5.23	5.82	2.85	4.55	3.78				
HEMBA1004656	3.49	2.49	3.49	3.55	3.42	3.65	2.19	3.03	2				
HEMBA1004657	23.62	14.49	14.4	48.51	47.67	43.85	51.21	56.08	58.34	**	+	**	+
HEMBA1004666	1.8	1.42	1.03	2.78	2.47	2.72	1.97	2.35	2.06	**	+	*	+
HEMBA1004669	5.4	3.16	2.99	6.16	6.23	6.59	2.94	2.65	2.66	*	+		
HEMBA1004678	4.37	2.24	2	5.27	6.01	4.17	2.94	3.39	4.41				
HEMBA1004672	5.55	2.84	2.98	5.68	8.28	8.14	3.49	6.01	3.36	*	+		
HEMBA1004689	43.34	14.93	30.58	21.98	24.65	26.05	13.05	14.68	11.72				
HEMBA1004690	4.61	2.61	2.69	2.94	2.18	2.84	1.97	4.01	2.41				
HEMBA1004693	2.15	1.25	1.33	2.01	3.2	3.06	1.39	3.08	2				
HEMBA1004697	7.39	3.61	2.79	3.75	7.36	9.2	5.36	4.7	6.2				
HEMBA1004702	21.02	14.02	11.62	9.2	10.6	12.82	11.9	12.65	12.79				
HEMBA1004704	6.08	3.81	3.24	8.5	8.45	8.19	4.75	5.52	5.39	*	+		
HEMBA1004705	1.15	0.61	0.21	1.49	1.26	1.73	1.37	1.44	1.36				
HEMBA1004706	3.9	2.72	2.07	2.01	3.27	2.47	3.18	2.94	2.37				
HEMBA1004709	3.4	2.4	2.61	5.18	5.97	7.11	2.19	3.92	3.09	**	+		
HEMBA1004711	3.02	1.29	2.07	2.19	3.63	3.64	1.38	3.22	1.38				
HEMBA1004723	9.52	5.41	7.44	9.15	11.88	10.6	1.92	9.59	6.27				
HEMBA1004725	5.24	3.87	3.31	6.21	5.61	5.19	5.65	5.85	6.52		*	+	
HEMBA1004729	1.7	2.99	1.13	11.04	3.71	3.48	1.14	4.24	1.15				
HEMBA1004733	1.86	1.11	1.27	1.93	2.88	2.54	1.38	2.89	2.03	*	+		
HEMBA1004734	2.06	1.99	1.5	2.15	2.83	2.82	2.1	2.82	2.29				
HEMBA1004736	3.46	3.3	2.73	5.69	8.26	7.15	2.94	3.83	4.08	**	+		
HEMBA1004748	4.24	1.57	1.93	4.83	6.28	6.83	2.64	4.21	2.49	*	+		
HEMBA1004749	7.35	4.59	3.33	5.23	6.38	10.26	4.24	7.9	6.41				
HEMBA1004751	3.74	2.05	2.99	5.29	6.07	7.15	2.9	5.44	3.62	*	+		
HEMBA1004752	5.63	3.06	2.11	4.83	5.66	7.24	4.55	3.43	5.1				
HEMBA1004753	85.27	60.35	45.13	73.61	76.67	82.04	35.88	33.51	34.18				
HEMBA1004755	12.21	10.42	8.56	18.13	22.58	19.53	19.43	13.65	17	**	+	*	+
HEMBA1004756	1.98	0.4	0.9	1.17	2.37	1.88	1.4	2.86	2.34				
HEMBA1004758	3.05	2.33	2.23	5.05	4.15	4.14	2.36	3.68	2.72	**	+		
HEMBA1004763	2.53	2.54	2.42	3.64	3.57	2.87	2.52	4.35	2.6	*	+		
HEMBA1004768	0.63	0.57	0.48	2.03	2.91	1.86	1.11	2.85	0.94	**	+		
HEMBA1004770	1.17	0.28	1.04	3.43	2.96	3.94	2.26	2.05	1.62	**	+	*	+
HEMBA1004771	3.01	1.5	1.36	3.25	3.04	3.46	2.2	2.12	2.05				
HEMBA1004773	6.8	4.62	3.7	7.13	8.07	9.04	7.62	7.16	8.82	*	+		
HEMBA1004776	3.71	2.57	1.18	9.61	3.42	2.31	3.1	3.48	4.38				

22/180

【0663】

【表192】

1521

1522

HEMBA1004778	4.28	3.09	3.12	5.87	7.81	8.46	5.37	4.86	3.66	*	+		
HEMBA1004774	1.55	1.14	0.87	1.97	2.67	2.4	1.81	2.87	1.66	*	+		
HEMBA1004785	2.2	0.85	1.41	2.94	2.11	2.82	2.94	3.76	2.42		*	+	
HEMBA1004789	2.02	2.15	2.94	6	4.66	4.07	4.12	6.23	6.59	*	+	*	+
HEMBA1004793	1.94	0.91	1.99	4.74	2.62	2.39	1.99	2.85	2.46				
HEMBA1004797	3.34	1.51	1.57	3.19	4.14	4.19	3.42	2.94	4.14				
HEMBA1004803	1.73	1.53	0.52	3.19	3.28	3.24	3.3	2.11	2.68	**	+	*	+
HEMBA1004806	1.99	0.24	0.76	2.51	2.13	1.62	1.14	2.33	1.41				
HEMBA1004807	6.07	4.25	4.5	4.85	8.03	9.33	4.48	5.59	5.41				
HEMBA1004816	3.49	2.36	1.89	3.34	3.8	3.31	2.37	4.02	1.69				
HEMBA1004820	1.49	1.14	1.32	2.51	2.88	2.8	1.5	4.47	1.86	**	+		
HEMBA1004833	7.98	3.57	4.1	7.09	8.03	7.72	4.99	7.63	6.59				
HEMBA1004847	6.33	4.11	5.21	8.38	7.16	8.48	4.35	8.93	6.34	*	+		
HEMBA1004850	3.92	2.57	2.41	5.26	3.09	3.63	3.54	3.4	5.81				
HEMBA1004863	4.26	1.79	2.07	6.34	5.16	5.37	2.36	2.91	5.42	*	+		
HEMBA1004864	8.29	3.32	3.08	5.48	7.27	7.94	4.75	3.71	4.59				
HEMBA1004865	1.92	1.18	0.62	2.11	6.7	3.86	2.14	1.94	1.68				
HEMBA1004880	4.54	3.09	3.36	6.03	7.12	7.25	3.5	4.7	4.49	**	+		
HEMBA1004882	5.35	4.05	3.06	4.2	4.72	3.45	2.62	4.51	3.09				
HEMBA1004883	1.17	0.68	0.57	1.14	0.82	0.86	0.53	1	0.47				
HEMBA1004899	3.26	2.08	1.7	3.09	2.94	3.37	2.23	2.83	5.72				
HEMBA1004900	1.39	1.1	0.25	1.7	1.35	1.1	1.57	1.47	1.61				
HEMBA1004902	6.14	4.05	3.74	6.91	8	7.96	4.94	4.32	3.82	*	+		
HEMBA1004918	4.98	2.15	2.73	5.38	6.39	6.51	3.65	3	3.79	*	+		
HEMBA1004923	1.88	1.64	1.69	3.18	2.96	3.02	2.23	2.61	2.53	**	+	**	+
HEMBA1004929	2.42	1.04	1.11	2.68	2.08	2.3	2.43	1.05	1.27				
HEMBA1004930	5.54	5.02	5.16	8.04	11.27	11.38	5.24	6.2	5.58	*	+		
HEMBA1004933	2.24	1.54	1.06	2	2.4	2.08	1.19	1.47	2.06				
HEMBA1004934	0.55	0.77	0.07	1.15	0.99	1.58	1.85	2.74	1.58	*	+	*	+
HEMBA1004937	6.5	2.53	3.22	3.69	3.97	5.19	4.16	4.2	3.69				
HEMBA1004943	6.44	2.93	2.55	5.45	3.9	5.9	3.81	4.39	5.14				
HEMBA1004944	4.47	1.97	2.6	5.4	4.69	6.01	3.98	3.08	5.3	*	+		
HEMBA1004946	6.58	4.26	2.56	8.23	7.78	9.16	5.73	6.06	6.35	*	+		
HEMBA1004952	5.05	2.8	1.43	3.17	3.73	3.4	2.89	3.56	3.26				
HEMBA1004954	2.94	2.13	2.53	7.6	9.09	8.39	8.28	11.47	6.83	**	+	*	+
HEMBA1004956	1.7	0.98	0.85	2.16	2.35	1.65	2.19	1.65	0.68				
HEMBA1004960	4.22	1.35	1.83	3.33	4.35	3.89	3.18	2.33	2.62				
HEMBA1004971	2.85	2.08	2.33	3.11	3.19	2.48	4.48	3.31	3.12				
HEMBA1004972	7.97	3.44	5.28	7.05	7.91	7.94	4.91	4.41	4.71				
HEMBA1004973	4.05	2.96	1.6	4.3	3.46	4.03	3.1	2.58	3.76				
HEMBA1004977	14.24	10.04	6.48	10.74	14.23	17.72	5.8	5.62	5.43				
HEMBA1004978	3.63	3.21	1.82	4.34	4.05	5.53	3.79	4.18	2.53				
HEMBA1004980	2.51	2.43	1.78	4.29	5.14	5.81	2.73	3.03	2.97	**	+		
HEMBA1004982	1.4	0.95	0.55	1.1	1.4	2.11	0.94	2.12	0.81				
HEMBA1004983	1.7	1.5	1.07	1.2	1.85	1.29	1.7	1.13	1.37				
HEMBA1004993	4.75	4.53	4.44	5.51	5.64	4.6	3.99	3.69	4.53				
HEMBA1005004	4.11	3.34	2.48	4.8	3.41	3.91	3.87	2.23	2.59				
HEMBA1005008	5.55	2.4	3.38	3.53	5.55	4.97	3.01	4.08	3.22				
HEMBA1005009	10.15	9.95	7.66	7.94	11.36	8.06	4.21	5.06	5.68		**		
HEMBA1005019	6.33	2.93	3.49	4.34	5.65	6.43	4.58	6.41	4.98				
HEMBA1005031	5.34	2.42	3.36	5.76	5.02	6.07	3.05	4.52	3.35				
HEMBA1005032	7.09	2.85	5.15	5.79	6.77	7.31	4.12	4.52	4.02				
HEMBA1005035	13.29	11.61	9.27	16.88	22.86	20.98	13.14	12.08	14.81	*	+		
HEMBA1005036	9.37	4.9	6.57	4.71	7.3	8.39	7.97	8.49	9.16				
HEMBA1005039	2.56	2.26	1.97	3.46	4.91	5.18	2.9	3.33	3.77	*	+	*	+
HEMBA1005047	3.73	2.69	2.58	2.7	3.22	4.69	3.19	3.28	3.52				

23/180

【0664】

【表193】

1523

1524

HEMBA1005950	8.01	4.69	4.35	6.4	8.24	6.75	4.64	5.95	4.47						
HEMBA1005962	2.24	3.49	0.58	2.31	2.34	1.56	1.28	2.55	1.35						
HEMBA1005966	1.59	0.53	1.22	1.43	2.19	2.08	0.94	1.37	2.12						
HEMBA1005967	10.97	5.24	5.8	11.93	6.24	15.81	3.48	6.87	4.63						
HEMBA1005970	54.34	32.66	23.12	7.48	7.25	9.46	4.96	6.22	5.67						
HEMBA1005975	4.78	2.93	2.39	9.53	8.99	8.84	5.77	6.79	5.19	**	+	+			
HEMBA1005978	9.58	7.81	5.77	9.39	9.72	10.05	5.01	5.86	6.72						
HEMBA1005979	12.04	7.57	6.48	19.42	17.72	15.43	8.5	9.75	11.45	*					
HEMBA1005983	2.66	1.46	0.66	1.94	3.02	2.07	1.27	2.07	1.73						
HEMBA1005984	7.91	6.72	4.77	5.71	7.83	8.74	5.49	4.34	3.97						
HEMBA1005989	2.86	1.68	1.86	2.41	5.46	5.18	1.67	3.51	1.49						
HEMBA1005992	5.98	4.14	4.3	9.36	10.56	9.53	2.93	3.59	4.57	**	+				
HEMBA1005990	33.54	22.43	17.55	44.06	40.43	42.47	20.48	22.61	17.3	*					
HEMBA1005996	5.76	3.98	4.37	6.03	5.87	6.22	4.23	3.3	6.47						
HEMBA1005101	5.71	2.76	3.85	3.75	5.23	3.72	2.48	3.36	3.77						
HEMBA1005107	4.5	1.82	2.91	2.69	3.89	3.12	3.06	3.74	2.52						
HEMBA1005113	1.45	0.81	0.45	8.23	11.09	10.71	5.43	5.23	6.51	**	+	**	+		
HEMBA1005123	10.61	5.86	5.3	15.09	21.59	18.64	8.57	8.85	8.33	*					
HEMBA1005133	2.6	2.55	2.08	3.44	6.93	6.67	3.17	4.12	2.67	**	+				
HEMBA1005135	1.91	1.13	1.66	1.75	3.38	1.54	1.31	3.02	1.14						
HEMBA1005145	16.67	9.87	9.21	12.39	15.8	16.28	8.2	10.27	10.61						
HEMBA1005149	10.32	5.61	5.06	11.44	12.5	12.06	7.17	8.59	7.71	*	+				
HEMBA1005157	6.34	4.06	3.55	9.52	11.4	12.18	3.28	4.04	5.79	**	+				
HEMBA1005159	0.7	1.49	0.94	1.57	3.26	1.76	1.22	2.54	0.91						
HEMBA1005172	43.22	25.23	24.37	33.5	39.86	37.96	32.09	25.74	34.44						
HEMBA1005185	4.97	4.57	2.99	2.86	3.27	4.08	2.48	3.14	1.7						
HEMBA1005186	3.35	2.42	3.23	5.64	6.25	4.46	2.06	2.21	2.79	*	+				
HEMBA1005195	1.99	0.84	0.81	1.89	2.31	1.52	1.31	2.87	1.25						
HEMBA1005201	6.2	5.19	2.55	3.77	6.88	6.27	4.89	5.22	6.55						
HEMBA1005202	8.96	4.63	5.23	6.96	8.01	6.67	8.1	7.62	9.46						
HEMBA1005204	113.3	93.42	81.36	145.9	165	106.5	90.09	59.5	89.11						
HEMBA1005206	6.48	3.93	4.83	5.9	5.71	6.15	4.98	4.32	4.52						
HEMBA1005212	2.14	1.72	1.8	4.03	2.98	2.85	3.28	4.04	4.31	*	+	**	+		
HEMBA1005223	3.02	2.16	2.78	4.29	3.41	4.21	2.9	3.66	3.28	*	+				
HEMBA1005229	0.71	0.07	0.59	1.25	1.02	0.47	0.51	2.08	0.98						
HEMBA1005236	4.24	4.62	2.37	7.34	7.76	6.64	2.52	4.81	4.22	*	+				
HEMBA1005232	0.15	0.54	2.47	1.05	1.44	1.37	1.1	0.73	0.86	**	+	+	+		
HEMBA1005238	5.05	3.37	2.42	6.46	5.11	6.11	4.05	3.86	3.91						
HEMBA1005241	18.2	11.3	9.41	11.74	14.66	18	9.85	7.33	9.11						
HEMBA1005244	6.45	3.35	4.4	5.3	7.24	5.83	3.98	5	6.42						
HEMBA1005246	9.39	6.95	6.65	15.52	17.83	13.37	15.28	9.28	12.95	**	+				
HEMBA1005251	2.49	1.43	2.18	5.25	6.15	4.92	3.41	3.93	3.36	**	+	+	+		
HEMBA1005255	3.83	2.63	3.03	3.56	4.92	3.46	2.88	4.5	4.38						
HEMBA1005267	1.63	0.84	1.67	10.27	7.55	7.28	1.17	3.13	1.81	**	+				
HEMBA1005274	1.18	0.71	0.61	1.46	2.14	1.62	1.18	1.08	1.02	*	+				
HEMBA1005275	1.9	0.81	0.85	2.82	4.11	3.4	1.92	2.54	1.27	*	+				
HEMBA1005288	3.5	1.84	2.36	6.6	8.72	6.93	3.45	3.43	3.32	**	+				
HEMBA1005293	1.91	2.03	0.55	4.54	1.95	1.33	0.58	2.15	1.2						
HEMBA1005296	401.9	314.1	296.6	377.8	403.7	432.4	228.1	207.4	230						
HEMBA1005301	1.98	0.74	1.57	2.67	1.35	1.8	1.62	2.6	1.33						
HEMBA1005304	4.1	2.37	2.93	8.99	8.69	9.63	4.7	6.26	5.96	**	+	+	+		
HEMBA1005305	2.8	1.25	1.81	4.03	4.81	4.66	2.57	4.35	2.78	**	+				
HEMBA1005311	2.04	1.03	1.55	2.81	2.74	3.45	0.88	2.3	2.96	*	+				
HEMBA1005313	6.91	3.99	3.19	6.31	4.42	4.78	4.14	4.74	6.56						
HEMBA1005314	0.55	0.27	0.2	1.02	1.14	0.89	1.2	4	1.03	**	+				
HEMBA1005315	4.12	1.27	1.36	3	4.13	3.44	3.48	2.53	2.28						

1525

1526

HEMBA1005317	1.33	0.36	0.19	4.23	3.8	4.6	1.25	1.46	1.92	**	+		
HEMBA1005318	1.08	0.85	0.59	0.97	1.89	1.29	1.43	1.82	1.13				
HEMBA1005324	3.04	2.4	1.83	6.59	7.62	7.75	5.26	6.51	7.55	**	+	**	+
HEMBA1005331	0.95	1.56	1.2	1.7	1.65	2.13	0.66	2.53	0.91				
HEMBA1005337	2.8	1.37	1.32	2.67	3.1	2.37	2.01	2.34	2.18				
HEMBA1005338	4.38	1.6	2.45	4.11	1.92	3.95	3.55	3.33	3.26				
HEMBA1005344	22.24	11.71	11.54	14.09	14.09	14.6	12.65	14.29	14.22				
HEMBA1005353	6.55	4.18	3.72	6.77	13.54	9.81	6.95	6.75	7.1				
HEMBA1005359	7.54	5.12	6.63	11.85	12.2	12.76	7.38	8.41	9.39	**	+		
HEMBA1005363	9.18	7.14	7.14	5.77	8.95	8.4	3.09	3.31	2.6			**	-
HEMBA1005364	0.69	1.26	0.41	1.96	2.44	1.02	1.19	1.6	1.68				
HEMBA1005367	3.22	2.29	1.05	4.88	6.98	6.68	5.63	8	6.43	*	+	**	+
HEMBA1005372	2.2	0.98	0.77	1.74	3.83	3.08	4.16	2.78	2.66			*	+
HEMBA1005374	6.99	3.71	3.35	12.54	10.52	8.75	6.1	6.58	7.22	*	+		
HEMBA1005379	1.84	1.63	1.2	1.2	1.49	2.65	1.75	1.09	1.97				
HEMBA1005382	7.86	4.67	5.2	10.89	7.83	8.14	5.58	6.98	6.52				
HEMBA1005384	4.42	2.21	2.13	6.74	6.14	5.84	4.87	4.21	4.01	*	+		
HEMBA1005386	6.04	3.65	3.38	6.45	5.92	6.1	5.2	4.67	5.78				
HEMBA1005389	5.36	3.94	2.77	5.75	6.88	6.02	2.6	5.56	3.66				
HEMBA1005394	6.27	3.67	3.58	3.93	4.59	4.22	2.21	4.81	3.15				
HEMBA1005403	11.32	8.45	6.9	16.3	23.03	11.57	16.03	13.06	13.2			*	+
HEMBA1005408	4.6	4.51	2.17	5.61	4.87	4.3	5.51	3.2	4.27				
HEMBA1005410	1.48	1.46	0.96	2.22	1.83	2.32	3.82	2.31	2.31	*	+	*	+
HEMBA1005411	3.32	2.25	1.72	8.56	7.19	8.45	4.84	3.85	4.74	**	+	*	+
HEMBA1005423	4.84	2.65	2.83	7.04	5.69	5.75	3.26	4.32	3.64	*	+		
HEMBA1005426	1.66	0.94	1.03	2.84	2.24	2.73	1.74	2.79	1.34	**	+		
HEMBA1005427	18.06	13.04	14.1	24.89	25.18	27.94	11.55	18.31	15.99	**	+		
HEMBA1005430	3.16	1.5	2.13	1.75	2.9	3.77	2.43	3.98	2.23				
HEMBA1005438	4.91	3.54	3.44	5.97	8.41	5.02	5.97	4.67	6.58				
HEMBA1005443	11.24	11.79	6.21	19.21	19.58	15.66	17.03	13.17	10.83	*	+		
HEMBA1005447	3.13	3.2	1.74	4.18	4.12	4.68	2.92	2.36	2.86	*	+		
HEMBA1005449	4.87	2.92	3.15	2.75	4.63	3.51	2.81	3.38	5.99				
HEMBA1005452	8.28	4.39	4.04	3.56	7.29	6.13	4.29	5.16	4.62				
HEMBA1005454	6.03	4.13	3.77	3.63	4.31	5.36	2.84	5.74	3.18				
HEMBA1005468	8.63	4.08	5.4	8.19	9.21	9.17	5.46	7.18	6.46				
HEMBA1005469	7.04	4.49	4.09	8.04	6.87	9.35	3.55	5.47	4.98				
HEMBA1005472	4.58	4.13	2.33	5.09	7.14	6.31	4.57	3.72	5.09				
HEMBA1005474	7.99	6.35	8.53	12.45	17.71	14.57	6.84	6.03	7.86	*	+		
HEMBA1005475	27.06	16.75	12.04	21.27	20.2	24.59	14.7	11.72	14.55				
HEMBA1005489	4.67	3.91	3.31	12.33	12.95	12.78	5.02	3.73	4.43	**	+		
HEMBA1005497	1.7	0.87	0.7	1.28	2.32	1.65	1.49	1.73	0.9				
HEMBA1005508	6.11	2.66	2.28	6.01	8.49	7.76	2.99	5.44	4.21				
HEMBA1005506	3.91	0.96	0.87	1.02	1.78	1.61	1.14	3.14	1.21				
HEMBA1005508	3	1.68	2.62	3.65	3.78	4.7	1.31	2.01	2	*	+		
HEMBA1005511	6.78	4.02	3.71	12.46	10.15	10.8	6.67	5.32	7.37	**	+		
HEMBA1005513	9.39	4.07	4.88	7.16	6.69	8.41	5.04	6.55	4.92				
HEMBA1005517	4.77	2.9	3.52	2.59	3.48	4.27	1.92	3.32	2.46				
HEMBA1005518	6.02	2.95	2.57	4.55	4.62	5.87	5.99	3.9	5.53				
HEMBA1005528	11.23	5.82	6.06	14.5	18.42	18.84	7.99	9.11	9.67	*	+		
HEMBA1005522	4.58	1.74	1.96	2.63	3.4	3.05	1.78	3.18	2.26				
HEMBA1005526	4	2.06	4.25	8.46	10.15	10.58	3.26	5.34	5.19	**	+		
HEMBA1005528	14.83	10.54	9.95	13.03	18.88	16.21	6.65	7.84	6.97			*	-
HEMBA1005536	5.44	2.29	3.17	4.84	6.25	8.18	4.67	4.21	3.84				
HEMBA1005538	4.71	2.93	2.46	83.2	102.3	97.16	227.3	162.3	210.6	**	+	**	+
HEMBA1005539	7.02	4.61	3.84	4.34	5.62	5.7	5.14	4.99	5.58				
HEMBA1005545	4.05	4.59	3.18	3.31	5.22	4.49	4.33	4.46	3.97				

25/180

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【表195】

1527

1528

HEMBA1005448	2.54	2.07	2.02	3.97	6.52	4.14	3.37	3.9	3.32	*	+	**	+
HEMBA1005452	9.98	4.38	5.49	14.16	16.16	16.24	6.88	9.1	7.91	**	+		
HEMBA1005458	5.62	4.78	4.01	4.12	4.94	4.94	2.89	4.54	2.98				
HEMBA1005468	4.56	2.35	2.64	4.41	6.84	7.67	2.66	3.77	3.75				
HEMBA1005470	22.81	14.72	12.89	3.4	5.87	4.67	2.86	3.28	4.18	*	-	*	-
HEMBA1005476	3.57	2.9	1.76	5.63	4.9	6.27	3.31	4.43	3.65	*	+		
HEMBA1005477	3.28	1.8	1.85	2.52	3.76	3.29	1.78	2.45	2.1				
HEMBA1005481	6.44	3.47	3.35	11.86	10.8	9.38	9.31	8.35	7.77	**	+	*	+
HEMBA1005483	3.79	2.19	1.67	4.94	4.83	5.37	3.11	3.69	2.69	*	+		
HEMBA1005483	2.18	2.16	1.54	2.99	3.77	4.66	2.3	2.75	1.62	*	+		
HEMBA1005488	3.6	2.49	3.31	8.28	7.89	9.86	3.63	5.17	4.67	**	+		
HEMBA1005493	3.44	3.2	2.65	4.18	6.03	3.87	2.97	3.28	2.95				
HEMBA1005495	2.58	2.31	1.83	3.46	4.98	5.89	3.2	2.15	3.87	*	+		
HEMBA1005497	13.38	9.58	8.44	10.53	12.2	11.02	8.53	9.47	8.93				
HEMBA1005498	12.27	7.53	6.44	5.89	7.11	6.3	8.22	8.78	11.95				
HEMBA1005499	5.25	3.66	3.27	10.52	11.83	10.56	4.85	5.36	3.71	**	+		
HEMBA1005516	5.15	3.24	2.69	6.77	7.27	7.69	4.76	5.32	4.74	*	+		
HEMBA1005521	5.71	4.59	4.34	4.48	5.05	3.45	2.83	4.75	2.83				
HEMBA1005527	4.83	2.61	2.82	6.51	8.02	6.48	3.29	4.97	4.83	*	+		
HEMBA1005528	5.64	3.83	3.44	12.81	11.82	14.97	10.64	9.94	13.34	**	+	**	+
HEMBA1005531	2.21	1.39	0.65	2.83	4.04	3.15	5.61	3.11	3.88	*	+	*	+
HEMBA1005532	11.01	3.49	3.42	8.83	9.02	7.82	5.06	4.35	5.44				
HEMBA1005534	6.35	2.76	2.05	5.36	8.63	6.5	4.98	5	6.87				
HEMBA1005562	1.07	1.53	1.02	2.26	2.43	2.33	2.04	1.73	1.38	**	+		
HEMBA1005566	4.52	3.82	4.32	9.91	8.09	7.3	6.48	6.28	5.06	**	+	*	+
HEMBA1005570	2.29	2.27	1.9	7.3	6.51	7	3.1	7.04	3.71	**	+		
HEMBA1005571	3.97	1.07	3.6	3.68	3.22	2.26	4.53	6.9	3.6				
HEMBA1005579	4.26	2.11	3.13	6.55	7.51	6.35	2.51	4.92	3.8	**	+		
HEMBA1005580	6.79	3.09	2.88	6.98	9.15	8.11	7.19	3.45	6.54				
HEMBA1005583	5.15	2.24	2.86	3.16	3.75	6.06	3.75	2.67	3.13				
HEMBA1005598	6.46	4.64	3.65	6.51	6.49	8.04	4.48	5.97	6.27				
HEMBA1005599	2.04	1.37	1.03	2.33	2.8	2.44	1.39	3.16	0.93	*	+		
HEMBA1005703	1.57	1.14	0.53	2.63	1.8	1.22	0.95	3.02	1.71				
HEMBA1005705	4.78	2.62	3.65	8.55	5.59	7.85	3.94	5.46	2.65	*	+		
HEMBA1005712	1.7	0.73	0.42	2.78	2.29	2.36	1.03	2.79	1.13	*	+		
HEMBA1005717	1.99	1.9	1.57	4.59	18.53	4.07	1.65	3.65	2.24				
HEMBA1005718	12.46	6.17	5.4	10.4	11.53	8.97	6.74	7.19	8.25				
HEMBA1005721	13.4	8.95	6.41	11.18	12.64	11.59	11.3	10.89	13.73				
HEMBA1005723	11.88	7.25	5.73	15.89	16.63	13.24	10.07	13.96	12.55	*	+		
HEMBA1005724	4.23	1.39	1.12	1.47	3.11	2.3	1.44	1.83	2.83				
HEMBA1005732	4.64	3.73	2.82	4.17	4.78	5.5	3.41	2.84	3.27				
HEMBA1005737	2.11	1.17	0.89	1.64	1.86	1.55	2.37	1.99	3.73				
HEMBA1005742	2.91	1.85	1.65	20.12	22.7	20.93	10.11	6.75	7.19	**	+	**	+
HEMBA1005744	3.55	2.22	2.55	2.88	5.21	3.91	2.28	2.67	1.49				
HEMBA1005747	6.73	2.98	3.61	4.2	6.34	4.06	4.88	4.78	5.21				
HEMBA1005749	16	15.05	7.61	16.72	17.56	14.78	13.73	10.17	19.02				
HEMBA1005753	1.55	1.38	0.58	2.76	3.45	1.74	2.11	2.82	2.29			*	+
HEMBA1005760	6.22	4.23	3.01	5.27	5.19	5.24	4.36	3.24	4.73				
HEMBA1005763	5.47	4.02	4.47	8.82	8.58	6.98	4.72	5.79	3.58	**	+		
HEMBA1005766	6.49	3.72	3.07	6.86	5.34	6.17	4.5	5.2	3.85				
HEMBA1005788	5.24	3.72	3.56	7.77	10.48	12.03	5.65	6.93	5.8	*	+	*	+
HEMBA1005795	2.44	2.1	2.01	3.69	3.63	2.88	1.69	3.18	1.68	*	+		
HEMBA1005809	23.36	22	11.6	14.58	20.18	18.5	16.89	18.97	9.81				
HEMBA1005813	3.44	3.52	2.49	3.52	4.47	4.04	2.83	4.45	3.63				
HEMBA1005815	6.13	3.52	2.7	5.29	7.35	4.96	4.74	5.46	7.01				
HEMBA1005822	4.2	1.96	2.92	8.67	7.02	9.4	4.99	3.69	6.16	**	+		

26/180

1529

1530

HEMBA1005829	7.71	4.11	4.16	9.68	9.82	10.63	5.68	6.05	6.18	*	+		
HEMBA1005833	5.58	4.05	3.69	5.07	5.16	5.6	4.09	4.46	5.21				
HEMBA1005834	6.55	4.34	5.21	12.06	12.18	15.25	4.16	7.19	5.66	**	+		
HEMBA1005844	55.19	32.63	42.62	52.31	50.88	44.4	13.71	22.39	16.54		*	-	
HEMBA1005852	14.32	7.35	8.88	11.42	13.87	12.28	12.12	9.6	10.71				
HEMBA1005853	4.46	3.87	2.7	5.48	7.15	7.24	6.76	3.1	4.03	*	+		
HEMBA1005878	10.9	9.31	6.82	15.29	18.75	18.35	11.26	9.02	9.91	**	+		
HEMBA1005883	2.8	3.02	2.09	2.99	4.75	3.12	3.03	3.43	2.83				
HEMBA1005894	1.78	1.18	0.5	2.41	2.22	1.91	2.16	1.93	1.73				
HEMBA1005891	1.55	1.14	0.52	2.25	4.37	4.09	2.08	2.69	1.69	*	+		
HEMBA1005894	3.43	2.12	2.97	5.44	5.86	5.44	2.54	4.52	2.77	**	+		
HEMBA1005898	16.67	8.8	11.51	11.61	18.53	21.97	6.97	12.21	8.22				
HEMBA1005902	4.41	3.46	2.53	2.97	3.31	3.57	3.63	4.8	4.43				
HEMBA1005907	1.14	1	0.72	1.39	1.9	1.41	1.83	2.17	1.38		*	+	
HEMBA1005909	0.96	0.99	0.06	0.74	1.52	0.83	1.8	0.82	0.95				
HEMBA1005911	5.56	3.24	3.54	5.59	8.12	8.18	4.97	3.97	5.62	*	+		
HEMBA1005912	6.61	6.28	5.64	8.63	10.33	8.51	7.27	7.15	4.9	*	+	*	+
HEMBA1005913	3.32	1.87	2.67	4.85	5.83	5.39	4.23	6.09	5.19	**	+	*	+
HEMBA1005921	5.08	3.6	4.07	7.96	11.09	11.08	3.93	6.12	4.64	**	+		
HEMBA1005923	9.29	4.86	8.75	10.31	11.79	14.59	5.42	7.95	6.59				
HEMBA1005929	9.26	6.15	5.27	8.35	12.25	12.51	8.91	7.98	6.88				
HEMBA1005931	13.37	8.03	6.05	13.2	15.89	16.14	10.01	9.04	10.17				
HEMBA1005934	11.83	7.65	6.91	11.53	21.92	13.8	6.94	9.42	10.1				
HEMBA1005945	9.41	6.42	4.64	6.1	7.01	8.67	8.01	6.77	7.06				
HEMBA1005962	2.52	1.69	1.85	2.52	2.44	3.11	1.69	3.18	2.61				
HEMBA1005963	1.58	1.29	0.83	2.22	2.32	1.65	0.75	2.23	1.58				
HEMBA1005990	53.63	37.05	35.87	22.88	28.11	30.49	25.75	38.21	38.5				
HEMBA1005991	4.36	2.88	2.52	7.83	8.53	8.07	3.66	3.18	4.37	**	+		
HEMBA1005999	7.25	4.04	3.51	7.81	9.22	8.54	5.71	6.17	5.07	*	+		
HEMBA1006002	4.03	2.6	1.83	2.32	2.41	2.99	3.56	4.2	3.68				
HEMBA1006005	3.58	3.7	2.47	1.41	2.98	2.78	2.19	3.32	3.16				
HEMBA1006011	28.82	13.22	19.62	6.69	8.42	8.26	9.43	7.34	8.25	*	-		
HEMBA1006013	4.9	3.69	2.44	2.82	2.64	2.69	3.14	3.46	2.63				
HEMBA1006016	5.42	2.01	3.02	4.73	5.78	5.82	3.09	4.11	3.71				
HEMBA1006019	4.75	3.24	2.19	2.66	6.4	5.83	2.01	3.58	3.27				
HEMBA1006021	5.17	2.64	3.76	13.9	20.33	23.22	9.49	12.71	9.39	**	+	**	+
HEMBA1006022	6.7	7.43	3.24	7.5	7.39	6.93	5.83	6.01	8.3				
HEMBA1006031	4.39	5.2	2.1	3.55	7.12	4.25	2.82	4.39	3.34				
HEMBA1006035	3.57	1.83	2.1	2.68	3.31	3.32	3.52	3.36	3.1				
HEMBA1006036	11.47	5.72	5.91	13.84	22.61	19.36	7.96	7.38	10.66	*	+		
HEMBA1006042	5.24	3.69	2.84	6.48	8.01	7.56	4.36	7.77	4.18	*	+		
HEMBA1006044	1.69	0.79	0.7	2	1.1	1.58	0.9	2.05	1.25				
HEMBA1006045	4.3	3.06	2.36	5.33	6.87	5.75	4.69	7.34	3.91	*	+		
HEMBA1006048	5.42	3.01	4.33	5.37	6.23	4.19	3.1	3.81	2.5				
HEMBA1006053	5.79	4.06	2.48	4.5	6.49	3.55	3.66	3.74	4.34				
HEMBA1006055	1.82	1.84	1.28	1.8	2.36	2.19	1.75	2.52	2.48				
HEMBA1006058	4.72	2.18	2.21	2.56	3.95	3.04	3.54	3.28	3.39				
HEMBA1006063	15.52	11.99	10.03	16.08	16.03	13.94	13.46	9.12	10.83				
HEMBA1006067	1.98	1.55	1.25	1.72	2.65	1.7	2.65	2.72	2.71		**	+	
HEMBA1006081	3.98	3.25	2.94	3.52	4.19	3.86	2.74	3.6	2.18				
HEMBA1006089	10.88	7.08	9.01	8.5	7.48	9.62	5.26	4.65	6.13	*	-		
HEMBA1006090	2.72	1.74	2.31	2.48	4.09	2.53	1.71	3.25	2.66				
HEMBA1006091	8.41	4.97	5.38	6.67	13.08	9.53	6.54	5.82	7.45				
HEMBA1006093	4.66	3.46	1.8	4.22	4.68	5.91	4.02	4.39	5.59				
HEMBA1006099	8.2	2.83	3.57	7.25	7.27	6.62	8.7	6.8	7.75				
HEMBA1006100	4.94	3.58	3.48	6.75	7.94	7.84	5.43	4.52	4.63	**	+		

27/180

【0668】

【表197】

1532

HEMBA1006108	5.03	2.45	2.82	5.62	4.96	3.72	3.28	3.95	3.28								
HEMBA1006114	5.25	4.63	5.08	7.3	10.42	7.17	4.76	5.44	5.87	*	+						
HEMBA1006121	6.32	2.33	4.31	5.84	6.44	7.33	4.17	6.55	4.7								
HEMBA1006124	3.12	2.28	2.5	3.3	4.9	2.3	1.89	3.9	2.53								
HEMBA1006125	10.14	8.44	4.52	7.52	17.2	16.18	9.52	10.87	14.31								
HEMBA1006130	2.62	2.68	2.39	2.72	3.08	4.43	3.7	4.3	4.1			**	+				
HEMBA1006138	7.26	4.73	3.72	9.3	11.39	10.14	5.49	5.98	7.37	*	+						
HEMBA1006142	6.22	3.63	4.24	7.33	10.18	10.72	6.57	6.34	6.19	*							
HEMBA1006150	16.28	10.88	9.66	15.57	15.3	13.33	6.57	7.84	7.68								
HEMBA1006151	8.94	6.23	8.3	9.44	9.41	9.8	14.8	13.36	17.11			**	+				
HEMBA1006156	4.31	2.12	3.11	2.99	2.19	2.62	2.75	4.44	3.92								
HEMBA1006158	1.99	2.23	1	5.52	2.28	1.62	0.79	3.02	2.04								
HEMBA1006164	7.82	6.53	4.48	10.95	14.83	12	6.46	6.96	7.98	*	+						
HEMBA1006171	3.78	1.96	1.78	2.93	3.7	4.2	6.07	5.07	5.46					+			
HEMBA1006173	3.13	1.34	2.45	2.99	4.82	4.35	2.87	4.45	2.71								
HEMBA1006176	17.29	15.19	12.08	17.22	16.16	22.1	76.2	63.22	78.98			**	+				
HEMBA1006182	2.42	1.06	1.52	2.8	3.22	2.43	1.16	3.5	1.94								
HEMBA1006197	6.41	5.46	4.82	12.92	9.66	9.7	4.32	5.89	5.15			**	+				
HEMBA1006198	9.58	7.2	6.52	9.4	9.55	10.32	5.65	8.56	6.79								
HEMBA1006213	2.56	0.9	1.99	3.02	4.19	4.18	1.76	2.58	3.01	*	+						
HEMBA1006217	23.61	12.95	14.09	28.71	29.21	22.65	54.8	57.77	74.75			**	+				
HEMBA1006226	45.81	48.81	55.06	71.05	67.87	69.04	34.7	30.76	48.77	**	+						
HEMBA1006235	2.69	1.66	2.93	2.89	2.63	3.42	3.26	2	2.73								
HEMBA1006248	4.57	1.66	2.14	4.47	3.25	4.51	3.57	3.35	2.98								
HEMBA1006251	7.31	5.13	5.62	8.77	8.46	10.33	8.03	7.68	7.92	*	+			+			
HEMBA1006252	2.83	2.65	0.78	1.86	3.33	3.7	2.51	1.94	2.08								
HEMBA1006253	5.52	3.08	3.71	4.06	4.47	4.75	2.99	2.68	1.89								
HEMBA1006259	4.17	1.88	2.86	4.37	4.68	6.45	2.66	2.31	3.69								
HEMBA1006261	6.4	3.25	3.39	6.02	5.83	6.2	5.45	3.63	10.61								
HEMBA1006266	3.66	2.08	1.88	4.46	4.9	5.18	2.58	2.36	4.27	*	+						
HEMBA1006271	7.71	2.93	4.51	11.62	12.09	12.3	7.07	3.33	10.91	**	+						
HEMBA1006272	2.81	1.63	1	2.86	2.92	3.49	2.16	1.96	2.4								
HEMBA1006273	3.39	2.09	3.07	4.81	3.79	4.4	5.32	3.06	3.91								
HEMBA1006276	2.93	1.9	3.24	3.4	4.55	3.76	2.55	1.66	2.29								
HEMBA1006278	1.93	1.63	1.33	4.06	4.19	3.8	2.43	1.58	2.09	**	+						
HEMBA1006283	7.35	3.25	3.5	4.82	5.8	5.93	4.92	3.12	4.11								
HEMBA1006284	3.83	2.26	2.04	5.58	2.8	4.34	3.15	2.33	3.82								
HEMBA1006291	4.96	1.36	1.34	4.1	2.68	4.41	3.86	3.13	3.18								
HEMBA1006292	2.77	2.02	1.73	2.32	2.22	1.89	2.26	1.67	2.38								
HEMBA1006293	3.02	0.92	1.7	1.99	1.76	2.36	1.34	1.85	1.56								
HEMBA1006299	3.49	2.22	1.51	13.99	12.83	16.92	7.99	7.28	10.15	**	+	**	+				
HEMBA1006309	5.39	3.08	3.38	5.38	6.85	7.74	3.06	4.11	4.45								
HEMBA1006310	3.7	2.35	2.24	5.29	3.06	3.56	2.99	4.56	4.32								
HEMBA1006311	8.15	4.04	4.72	3.8	4.97	7.43	4.03	5.26	5.64								
HEMBA1006313	2.58	0.57	1	1.55	1.73	1.85	2.63	1.09	1.3								
HEMBA1006316	2.99	1.66	1.44	1.74	2.62	2.14	2.59	1.79	1.84								
HEMBA1006324	4.68	2.1	1.68	6.39	5.95	6.83	4.37	3.72	3.95	*	+						
HEMBA1006334	2.26	1.44	1.07	1.93	1.34	1.2	1.12	1.33	1.91								
HEMBA1006335	10.13	6.95	5.67	4.72	4.51	6.4	10.88	11.65	14.09			*	+				
HEMBA1006344	4.43	2.82	4.27	2.97	8.14	7.72	4.65	6.26	4.98	**	+						
HEMBA1006347	5.25	2.13	2.64	4.75	3.92	6.02	3.83	3.69									
HEMBA1006349	6.07	2.73	2.89	4.44	4.96	6.67	4.94	4.8	4.22								
HEMBA1006352	3.21	2.07	2.2	4.23	3.53	3.79	3.57	2.77	2.87	*	+						
HEMBA1006357	9.36	4.79	5.03	14.77	14.32	14.23	7.21	5.46	6.81	**	+						
HEMBA1006358	4.06	2.27	1.93	3.39	4.53	4.11	2.36	2.11	2.2								
HEMBA1006359	11.9	9.22	8.59	18.27	21.46	21.84	9.68	9.92	7.59	**	+						

28/180

1533

1534

HEMBA1006360	7.98	4.95	5.62	5.47	3.56	4.4	1.94	2.36	2.5		*	-
HEMBA1006364	3.11	1.13	2.29	5.13	3.18	5.17	4.53	10.75	7.49		*	+
HEMBA1006377	9.83	4.08	4.81	9.68	6.73	11.12	5.12	6.15	6.04			
HEMBA1006380	8.33	2.76	3.16	7.63	7.47	9.64	4.68	4.63	5.81			
HEMBA1006387	27.84	15.11	15.63	23.73	22.47	28.24	18.48	12.8	18.34			
HEMBA1006388	9	3.81	3.21	10.12	10.82	10.06	5.78	6.06	7.86			
HEMBA1006390	10.59	5.3	6.11	5.74	8.16	9.4	6.45	5.63	5.84			
HEMBA1006391	5.9	2.52	2.93	4.19	2.98	3.66	3.7	3.39	4.92			
HEMBA1006398	1.24	0.85	0.78	1.46	2.48	2.33	1.32	1.57	1.25	*	+	
HEMBA1006405	6.46	2.31	3.39	3.97	5.97	7.86	4.98	4.43	6.01			
HEMBA1006410	10.66	4.34	6.26	48.24	9.18	6.95	5.67	6.99	5.47			
HEMBA1006416	7.58	3.75	4.83	11.17	11.6	18.4	5.86	5.53	5.52	**	+	
HEMBA1006418	4.85	2.81	2.36	4.42	4.54	5.46	2.95	3.19	4.26			
HEMBA1006419	8.31	4.08	4.44	13	13.16	12.95	7.56	6.59	6.8	**	+	
HEMBA1006421	2.57	1.36	2.21	4.58	3.93	3.93	2.69	2.86	2.95	**	+	
HEMBA1006424	1.92	1.1	0.54	1.6	1.43	2	1.13	1.24	1.46			
HEMBA1006426	6.91	3.24	3.97	14.78	13.77	12.89	6.5	5.72	7.38	**	+	
HEMBA1006430	4.14	1.54	1.15	3.22	4.8	4.46	2.25	2.55	3.21			
HEMBA1006435	3.24	1.25	2.86	4.15	5.58	5.24	2.63	2.65	2.43	*	+	
HEMBA1006445	5.47	3.56	1.09	4.34	6.2	5.79	5.24	5.14	9.95			
HEMBA1006446	2.47	0.4	0.6	1.78	0.97	2.17	2.61	1.77	0.98			
HEMBA1006456	9.3	7.18	5.88	27.97	39.33	36.06	25.26	23.55	25.96	**	+	+
HEMBA1006461	3.9	2.47	2.09	3.96	6.32	5.5	3.18	2.5	2.97			
HEMBA1006467	3.36	2.3	2.41	1.89	3.11	2.94	1.06	2.01	1.22		*	-
HEMBA1006470	3.32	2.6	1.74	4.73	4.89	6.17	2.71	2.99	2.35	*	+	
HEMBA1006471	2.77	2.01	2.5	2.54	4.17	4.09	1.83	2.8	1.93			
HEMBA1006474	3.4	0.88	1.69	1.95	2.26	1.5	0.73	1.98	1.64			
HEMBA1006476	7.63	2.81	3.49	7.03	6.55	10.28	5.71	6.01	8.9			
HEMBA1006482	53.61	36.99	43.8	47.46	64.27	63.44	24.67	21.43	26.34		*	-
HEMBA1006483	5.77	3.34	3.12	9.27	6.33	10.42	4.67	4.49	5.8	*	+	
HEMBA1006485	2.4	0.96	1.41	4.2	4.91	5.55	9.43	7.34	8.87	**	+	+
HEMBA1006486	22.07	14.47	14.17	13.5	21.65	20.32	9.55	5.18	8.79	*	-	
HEMBA1006489	2.84	0.31	0.23	0.63	1.22	0.91	1.3	1.95	0.72		*	-
HEMBA1006492	22.55	16.4	18.02	18.63	19.03	19.21	4.75	5.92	5.79		**	-
HEMBA1006494	1.6	0.13	1.42	1.49	1.22	1.56	0.94	0.97	0.8			
HEMBA1006497	4.42	2.46	1.3	2.7	3.58	4.13	3.19	2.22	2.93			
HEMBA1006501	6.77	2.17	3.41	4.37	3.72	6.05	2.94	2.94	4.13			
HEMBA1006502	14.3	11.26	8.46	15.96	17.52	16.95	15.96	11.43	17.31	*	+	
HEMBA1006507	3.4	0.72	1.23	5.85	4.08	5.84	2.92	3.88	4.16	*	+	
HEMBA1006517	4.63	2.62	2.31	5.72	6.14	5.82	3.68	4.27	4.71	*	+	
HEMBA1006521	3.02	1.72	1.98	2.24	2.77	2.97	3.41	3.45	2.36			
HEMBA1006523	6.54	3.38	7.96	6.72	7.42	7.81	5.9	6.56	6.87			
HEMBA1006530	1.54	0.77	2.01	2.93	1.8	2.4	1.35	1.69	1.44			
HEMBA1006535	2.61	2.15	0.64	3.13	3.63	3.67	2.05	1.48	2.17			
HEMBA1006536	5.93	3.85	4.16	6.52	8.47	8.22	4.62	4.48	4.48	*	+	
HEMBA1006540	4.27	2.17	1.9	4.22	2.42	3.65	2.42	2.05	2.05			
HEMBA1006544	1.52	0.67	1.46	2.15	3.36	3.6	2.21	2.99	2.6	*	+	+
HEMBA1006546	4.48	4.88	3.24	16.24	9.73	11.7	5.09	6.41	8.5	*	+	
HEMBA1006549	2.13	0.58	0.86	2.8	1.88	2.9	1.86	1.87	1.42			
HEMBA1006559	5.16	2.1	4.6	12.73	9.91	12.55	8.17	8.31	7.9	**	+	+
HEMBA1006561	2.22	0.76	1.85	3.22	2.69	2.84	1.63	3.25	2.16	*	+	
HEMBA1006566	1.5	1.62	0.13	0.8	1.28	0.97	1.14	1.33	0.88			
HEMBA1006569	4.26	2.46	1.96	4.02	5.76	5.28	3.58	2.64	3.6			
HEMBA1006572	1.59	0.24	0.54	0.56	0.89	1.09	1.05	1.04	1.21			
HEMBA1006579	2.51	1.31	1.43	2.63	2.93	3.26	6.37	6.01	6.92	*	+	+
HEMBA1006583	3.62	1.64	2.4	3.61	3.89	4.77	3.38	3.77	2.05			

29/180

【0670】

【表199】

1535

1536

HEMBA1006695	4.6	1.32	2.47	6.45	3.43	5.48	2.48	3.17	3.33				
HEMBA1006697	6.19	2.47	4	9.61	11.89	11.02	4.43	7.63	6.03	**	+		
HEMBA1006698	5.22	2.34	3.13	5.5	7.09	8.72	4.17	3.59	5.67	*			
HEMBA1006699	5.88	3.13	2.66	9.51	7.07	8.75	4.24	4.07	8.12	*			
HEMBA1006700	6.23	2.4	3.25	7.51	8.15	9.4	4.22	3.47	5.72	*	+		
HEMBA1006701	21.51	11.59	11.39	8.91	10.89	11.11	15.72	17.22	19.01				
HEMBA1006702	11.14	7.16	5.63	14.71	13.36	15.13	9.17	8.76	9.27	*	+		
HEMBA1006703	3.5	1.48	1.8	6.1	5.02	7.12	3.2	2.77	3.44	*			
HEMBA1006704	5.83	1.94	3.55	4.08	4.21	4.37	3.14	4	3.07				
HEMBA1006705	8.1	3.39	6.04	7.92	5.21	8.41	3.69	6.07	4.57				
HEMBA1006706	7.17	4.23	2.23	4.85	5.86	6.95	5.26	5.13	6.25				
HEMBA1006707	7.55	5.4	7.95	14.31	13.73	13.23	6.43	7.1	11.54	**	+		
HEMBA1006708	6.97	4.5	3.06	4.22	5.74	4.88	4.94	3.37	4.63				
HEMBA1006709	7.71	4.81	3.99	9.26	8.5	11.38	5.3	4.42	6.47	*	+		
HEMBA1006710	7.41	4.7	3.7	5.26	4.56	4.46	6.04	3.81	4.25				
HEMBA1006711	1.62	1.53	0.92	2.4	1.66	1.94	1.6	1.36	2.14				
HEMBA1006712	2.8	1.45	1.19	5.48	2.51	3.57	1.85	1.35	3.75				
HEMBA1006713	4.48	2.13	2.48	3.04	6.4	6.86	3.55	4.19	4.16				
HEMBA1006714	4.97	3.16	4.4	5.76	5.14	7.87	4.61	3.42	4.54				
HEMBA1006715	10.46	5.08	3.85	9.54	8.88	9.7	6.21	4.55	6.44				
HEMBA1006716	2.27	1.69	1.34	3.17	2.06	2.03	4.61	1.08	3.99				
HEMBA1006717	6.01	4.37	2.5	4.47	6.02	6.19	4.31	2.6	4.14				
HEMBA1006718	4.5	1.72	1.74	6.75	6.52	5.65	3.76	2.82	3.65	*	+		
HEMBA1006719	12.87	6.14	7.8	9.63	11.85	11.77	5.03	6.37	5.41				
HEMBA1006720	2.64	1.17	1.68	3.05	1.99	2.26	2.52	2.64	2.72				
HEMBA1006721	6.65	2.92	3.19	5.67	3.46	4.24	2.84	4.21	4.09				
HEMBA1006722	8.39	4.87	3.01	5.26	5	6.1	6.53	3.85	5.31				
HEMBA1006723	6.65	3.16	3.47	4.07	5.63	4.68	6.45	3.52	4.44				
HEMBA1006724	8.88	2.4	4.14	4.44	3.37	2.99	4.5	3.69	4.56				
HEMBA1006725	3.81	3.86	1.52	3.61	3.98	3.44	2.83	2.11	3.27				
HEMBA1006726	7.51	3.16	2.94	4.8	6.48	6.17	3.61	3.73	4.13				
HEMBA1006727	5.15	2.61	1.58	2.17	3.41	5.22	2.11	2.54	2.79				
HEMBA1006728	4.81	2.29	1.84	6.06	4.83	6.03	2.78	3.29	3.24				
HEMBA1006729	7.87	4.47	4.75	8.29	5.08	7.45	3.49	6.04	3.57				
HEMBA1006730	10.08	3.77	3.8	14.22	11.75	16.16	7.99	6.73	6.12	*	+		

1539

1540

HEMBA1007080	6.49	3.94	5.98	9.28	8.08	9.96	7.3	4.03	5.16*	+		
HEMBA1007084	6.15	4.73	3.3	7.53	11.43	12.96	4.4	6.54	5.84*	+		
HEMBA1007085	11.57	6.03	6.42	14.47	16.1	16.28	10.38	7.67	10.37*	+		
HEMBA1007087	8.74	3.56	3.06	9.07	6.88	8.89	6.19	4.83	6.54			
HEMBA1007099	4	1.06	1.76	3.78	3.23	3.34	2.04	2.11	2.69			
HEMBA1007095	70.95	56.95	68.33	67.92	65.09	77.99	58.54	73.69	65.61			
HEMBA1007101	8.13	4.48	3.34	8.09	6.98	8.85	8.22	8.26	10.62			
HEMBA1007104	5.96	2.99	2.91	5.64	4.63	4.55	3.31	2.95	4.74			
HEMBA1007106	14.7	8.59	9.92	8.69	10.08	8.52	4.85	5.28	6.28		*	
HEMBA1007112	2.54	1.7	2.5	2.24	3.01	4.69	1.68	1.95	2.11			
HEMBA1007113	6.43	3.26	3.02	9.57	10.18	12	5.01	4.74	6.51**	+		
HEMBA1007121	15.29	6.28	8.37	20.01	26.55	24.61	14	11.02	13.68*	+		
HEMBA1007129	4.97	2.15	2.01	4.97	3.24	4.17	3.35	2.12	2.39			
HEMBA1007147	3.38	3.64	3.3	8.42	7.76	8.81	5.26	3.94	5.49**	+		
HEMBA1007149	4.94	2.77	3.26	6.72	8.54	6.33	4.3	4.3	4.5*	+		
HEMBA1007151	8.13	3.85	3.81	6.44	7.95	11.22	3.61	4.33	3.79			
HEMBA1007172	7.56	3.48	4.13	7.44	5.05	7.28	4.12	5.13	4.04			
HEMBA1007174	5.89	2.49	3.67	3.93	3.83	5.88	2.72	4.89	2.91			
HEMBA1007176	9.03	5.34	6.92	9.78	8.83	9.47	8.52	8	9.63			
HEMBA1007178	32.35	18.88	15.14	19.06	21.65	17.59	9.08	8.47	7.89			
HEMBA1007185	10.22	4.41	3.64	8.36	9.55	9.52	7.01	9.23	10.25			
HEMBA1007186	5.79	5.42	2.99	5.38	6.38	4.34	5.53	4.63	4.28			
HEMBA1007194	10.77	5.25	6.27	6.05	8.58	8.52	4.54	4.6	4.86			
HEMBA1007200	4.17	3	2.87	3.85	3.81	6.07	2.25	5.2	3.91			
HEMBA1007203	7.33	3.38	4.4	6.6	5.9	7.76	3.26	5.38	5.31			
HEMBA1007206	5.36	1.62	4.58	8.87	7.23	9.37	4.17	4.51	4.13*	+		
HEMBA1007224	4.31	3.41	3.02	7.21	8.94	7.9	5.84	2.98	5.52**	+		
HEMBA1007226	8.11	2.53	3.92	5.1	4.65	5.56	3.57	3.89	3.99			
HEMBA1007240	8.19	3.25	3.14	6.63	3.95	5.13	4.82	3.47	3.83			
HEMBA1007241	2.29	1.82	2.1	4.38	3.16	4.31	2.93	3.05	2.77*	+	**	+
HEMBA1007242	3.53	1.89	1.63	1.79	3.06	2.59	1.23	2.17	2.42			
HEMBA1007243	5.49	1.9	2.36	5.15	4.7	4.56	2.5	2.45	3.07			
HEMBA1007281	3.85	1.52	2.26	3.21	2.8	3.04	1.54	2.44	1.83			
HEMBA1007286	2.11	1.7	2.58	4.85	3.63	4.4	1.15	2.23	2.07**	+		
HEMBA1007267	8.06	2.62	3.26	10.13	10.25	11.99	6.27	4.97	6.51*	+		
HEMBA1007273	2.76	1.75	1.08	1.92	1.89	2.71	1.52	1.78	1.02			
HEMBA1007279	2.55	1.22	1.16	1.3	3.65	2.92	1.5	1.89	2.13			
HEMBA1007281	2.07	1.07	0.43	1.29	1.21	1.04	1.02	1.25	1.12			
HEMBA1007283	6.62	2.63	3.23	3.75	3.81	4.38	2.73	2.29	3.75			
HEMBA1007288	3.75	1.29	2.66	5.75	6.28	6.21	1.78	2.89	4.17**	+		
HEMBA1007291	3.22	0.96	1.72	2.4	3.14	3.81	1.55	2.4	4.26			
HEMBA1007299	23.93	13.7	15.73	10.56	22.18	16.89	17.86	19.6	16.71			
HEMBA1007300	6.22	3.89	1.52	4.87	4.49	5.57	2.96	3.67	3.54			
HEMBA1007301	4.77	2.47	2.12	3.91	6.06	4.53	5.31	3.92	4.11			
HEMBA1007319	5.04	2.71	2.66	4.51	4.51	4.65	2.4	2.58	1.88			
HEMBA1007320	3.5	1.62	1.5	3	2.95	3.58	2.72	2.88	2.98			
HEMBA1007322	28.33	24.69	28.25	30.89	47.79	40.83	20.16	16.66	16.95	**	-	
HEMBA1007323	6.68	1.59	2.78	3.35	2.99	4.54	1.69	2.27	2.61			
HEMBA1007326	16.87	9.35	13.09	29.82	36.45	31.07	12.34	13.22	15.57**	+		
HEMBA1007327	6.34	3.6	4.38	10.61	13.22	12.6	4.55	6.34	5.25**	+		
HEMBA1007332	13.26	4.92	5.19	6.74	8.15	8.34	6.2	5.28	6.24			
HEMBA1007341	3.07	1.51	1.92	5.68	4.8	6.45	2.94	3.15	3.13**	+		
HEMBA1007342	3.54	1.8	1.84	3.52	2.33	2.69	2.06	2.55	1.53			
HEMBA1007347	6.86	4.49	4.81	9.76	12.67	13.86	6.9	5.92	8.38**	+		
HEMBA1007343	2.54	1.91	1.06	2.5	3.01	2.77	1.29	2.06	1.66			
HEMBA1008005	5.95	3.76	2.97	7.43	7.91	9.69	2.81	4.53	3.98*	+		

32/180

【0673】

【表202】

1541

1542

HEMBS1000006	6.33	3.99	3.35	9.32	9.9	11.83	4.69	4.68	5.58	**	+		
HEMBS1000018	9.18	4.31	7.12	14.89	18.9	20.93	7.15	7.95	8.65	**	+		
HEMBS1000024	8.61	5.92	3.83	12.18	15.58	14.42	6.22	5.52	8.3	**	+		
HEMBS1000025	7.18	1.68	2.62	3.76	5.35	5.09	4.63	4.5	5.11				
HEMBS1000030	5.99	4.74	5.88	11.95	12.01	10.44	5.68	5.83	6.43	**	+		
HEMBS1000036	5.65	4.09	3.36	4.79	4.59	7.69	4.76	4.78	5.5				
HEMBS1000037	6.62	4.31	5.17	7.83	6.16	9.26	6.18	5.41	5.32				
HEMBS1000039	3.3	1.35	2.08	5.56	6.46	6.46	3.88	3.39	2.84	**	+		
HEMBS1000044	8.31	2.86	3	8.94	8.97	9.22	3.67	5.53	3.74				
HEMBS1000048	4.16	1.72	3.61	5.69	6.15	8.14	3.51	4.43	3.25	*	+		
HEMBS1000050	5.5	1.49	1.55	3.76	8.59	5.41	2.51	2.18	3.82				
HEMBS1000054	5.55	2	2.33	9.07	6.03	8.7	7.15	3.88	5.66	*	+		
HEMBS1000055	24.4	16.2	17.8	18.24	19.34	22.83	9.69	8.54	9.54			*	-
HEMBS1000059	8.8	6.35	7.84	16.75	19.27	21.09	9.69	10.78	9.65	**	+	*	+
HEMBS1000072	9.51	4.64	5.32	12.83	10.68	11.19	7.97	7.6	5.64	*	+		
HEMBS1000081	3.87	1.35	1.85	5.08	5.24	4.46	3.77	3.99	4.68	*	+		
HEMBS1000083	4.74	2.08	3.56	8.88	6	6.36	3.2	5.07	6.07	*	+		
HEMBS1000089	3.6	2.1	3.13	10.31	7.12	8.77	3.62	4.07	4.02	**	+		
HEMBS1000094	10.03	4.21	5.44	7.27	9.1	10.43	5.68	3.83	7.07				
HEMBS1000097	2.21	1.8	1.66	3.6	3.78	2.43	2.31	1.65	1.94	*	+		
HEMBS1000099	6	2.44	5.07	9.23	13.61	11.37	6.57	5.71	7.13	*	+		
HEMBS1000103	11.08	5.29	6.37	9.34	10.14	10.72	4.69	6.24	4.67				
HEMBS1000106	6.42	4	5.39	8.37	6.27	6.82	6.5	5.47	4.71				
HEMBS1000113	2.17	2	1.61	3.56	3.45	3.36	1.25	3.37	2.9	**	+		
HEMBS1000119	4.55	2.45	4.15	5.3	3.89	4.98	2.17	5.09	5.65				
HEMBS1000133	36.74	19.87	32.19	17.43	2.43	25.47	18.03	19.17	26.05				
HEMBS1000134	8.1	5.02	4.94	5.99	6.85	11.63	3.4	5.64	6.33				
HEMBS1000136	4.52	2.17	1.45	2.82	2.31	2.54	3.01	2.62	4.93				
HEMBS1000141	5.34	2.26	2.68	7.34	8.23	8.82	4.82	3.93	6.2	*	+		
HEMBS1000144	4.28	3	3.58	12.18	6.95	9.35	4.11	4.95	6.86	*	+		
HEMBS1000147	3	2.36	0.48	3.68	2.83	3.66	1.75	1.4	2.8				
HEMBS1000152	4.26	2.59	2.98	3.85	2.52	3.5	2.62	3.23	3.16				
HEMBS1000154	3.63	1.65	1.97	5.05	4.98	5.15	2.28	3.46	4.23	*	+		
HEMBS1000155	3.1	2.14	2.06	3.13	4.38	4.5	2.17	2.09	2.04	*	+		
HEMBS1000173	11.42	5.05	6.29	19.61	16.74	17.56	10.24	8.45	9.62	**	+		
HEMBS1000175	3.73	1.02	1.8	5.42	5.67	6.02	2.9	2.66	4.4	*	+		
HEMBS1000176	5.82	2.57	3.52	6.79	7.3	6.93	5.44	4.12	6.38	*	+		
HEMBS1000198	2.93	1.33	0.9	2.24	0.81	1.87	1.77	0.77	1.87				
HEMBS1000208	3.02	2.41	1.68	3.5	2.31	3.21	2.28	1.81	1.61				
HEMBS1000209	4.47	2.11	2.26	5.05	5.4	5.79	2.1	3.16	2.24	*	+		
HEMBS1000212	4.74	2.38	2.45	3.32	2.97	6.08	1.78	3.81	2.18				
HEMBS1000215	12.22	6.74	7.81	16.21	19.51	21.21	10.04	11.3	10.31	*	+		
HEMBS1000217	18.97	9.31	7.7	15.35	13.44	12.33	8.45	8.5	11.37				
HEMBS1000218	7.28	3.65	4.15	11.14	12.99	13.65	6.32	5.71	6.71	**	+		
HEMBS1000226	9.75	5.82	3.67	9.36	7.18	7.09	5.55	5.63	5.67				
HEMBS1000230	2.5	1.54	1.56	3.16	2.41	2.47	1.66	1.86	2.28				
HEMBS1000240	2.54	1.04	1.59	2.21	2.34	2.83	1.25	2.23	1.86				
HEMBS1000244	3.34	2.45	3.05	3.32	3.23	3.4	1.54	2.3	1.85			*	-
HEMBS1000250	1.92	1.49	1.19	1.79	2.36	1.56	0.91	1.67	0.72				
HEMBS1000258	8.84	4.21	4.45	9.29	10.64	11.84	4.49	4.85	5.22	*	+		
HEMBS1000264	11.16	4.23	7.12	14.33	14.62	15.26	8.98	6.17	8.91	*	+		
HEMBS1000266	7.49	4.1	3.58	5.54	5.59	6.27	4.11	3.08	5.07				
HEMBS1000272	2.83	3.68	1.74	6.38	5.8	6.11	4.03	3.45	3.03	**	+		
HEMBS1000274	2.69	2.43	1.42	2.28	4.59	4.22	4.06	2.32	2.95				
HEMBS1000276	2.16	0.94	0.86	1.1	3.12	1.78	0.56	0.79	1.49				
HEMBS1000284	1.6	1.41	0.82	1.43	1.65	1.76	0.92	1.04	2.24				

33/180

【0674】

【表203】

1543

1544

HEMFB1000307	4.53	1.84	2.11	5.17	5.68	6.34	1.82	4.17	2.46	*	+		
HEMFB1000309	4.37	1.32	2.88	3.56	4.27	5.98	1.82	3.44	1.73				
HEMFB1000312	1.28	2.43	1.55	2.15	2.18	2.23	2	1.79	3.52				
HEMFB1000317	3.2	2.61	1.78	3.01	2.88	2.59	3.81	2.77	1.93				
HEMFB1000318	4.73	1.3	2.1	5.96	5.69	5.2	3.19	2.91	3.3				
HEMFB1000332	1.76	1.25	0.79	0.91	1.05	1.63	1.26	1.46	1.31				
HEMFB1000333	2.8	1.5	1.13	1.18	3.42	3.3	2.66	1.47	1.27				
HEMFB1000336	4.55	1.96	1.92	2.95	2.84	3.92	3.25	2.93	2.41				
HEMFB1000337	14.36	7.11	10.05	9.07	12	11.79	6.71	8.68	8.74				
HEMFB1000338	4.54	3.23	3.69	5.82	6.25	7.43	2.29	3.11	3.62	*	+		
HEMFB1000339	6.86	3.25	2.73	8.08	11.02	9.45	5.52	5.3	4.99	*	+		
HEMFB1000341	6.67	3.9	3.27	5.51	6.05	3.75	4.88	3.76	5.53				
HEMFB1000343	5.14	3.78	3.56	8.73	11.85	8.26	4.26	5.37	4.59	*	+		
HEMFB1000354	5.87	3.91	3.47	10.81	11.74	10.84	4.26	5.4	6.59	**	+		
HEMFB1000358	6.98	3.62	4.09	5.18	4.64	6.14	4.86	3.92	4.34				
HEMFB1000369	3.23	1.7	2.29	3.08	3.51	3.68	1.39	2.56	1.97				
HEMFB1000373	11.86	5.42	7.78	12.45	14.15	14.43	4.75	5.77	6.52				
HEMFB1000374	8.03	4.3	5.09	13.94	16.47	17.13	5.55	9.31	7.38	**	+		
HEMFB1000376	11.27	4.35	3.91	16.2	18.49	19.55	9.94	8.36	10.29	*	+		
HEMFB1000383	4.6	2.17	1.96	4.57	3.4	3.45	10.39	7.52	9.9		**	+	
HEMFB1000391	6.84	4.23	4.83	6	8.02	7.16	4.22	5.21	3.67				
HEMFB1000392	5.23	1.96	3.15	3.41	3.17	3.69	3.69	3.13	1.81				
HEMFB1000482	2.6	1.48	0.94	2.16	3.1	1.88	0.98	2.21	2.08				
HEMFB1000484	1.73	0.76	1.14	1.48	2.07	2.27	1.05	1.58	1.14				
HEMFB1000487	1.46	1.26	1.6	1.67	2.46	3.55	0.54	2.33	2.09				
HEMFB1000420	6.02	3.01	5.42	7.53	9.7	10.11	3.76	5.07	4.73	*	+		
HEMFB1000430	59.23	34.65	23.06	49.23	46.08	51.49	46.72	34.37	41.23				
HEMFB1000434	18.16	8.94	9.74	22.34	23.72	31.12	11.49	11.35	12.88	*	+		
HEMFB1000438	2.81	0.97	1.46	1.87	3.06	1.59	2.06	2.06	1.78				
HEMFB1000441	5.61	4.55	3.22	9.46	9.64	11.7	6.15	3.84	7.17	**	+		
HEMFB1000447	6.8	2.32	3.46	10.82	16.06	18.31	25.43	26.28	30.87	*	+	**	+
HEMFB1000449	1.31	0.73	0.5	2.05	2.12	2.41	1.36	2.6	1.7	**	+		
HEMFB1000453	8.09	6.85	8.91	11.38	10.07	15.36	7.99	10.3	12.98				
HEMFB1000455	2.58	3.4	2.03	3.63	4.91	3.97	1.67	3.24	1.52				
HEMFB1000472	7.59	4.06	3.3	4.71	4.91	6.8	5.17	4.42	5.06				
HEMFB1000480	9.8	3.69	3.57	8.18	11.17	10.77	5.35	5.7	6.17				
HEMFB1000486	7.07	2.27	3.48	8.16	9.71	10.13	5.36	5.39	6.03	*	+		
HEMFB1000487	2.41	1.44	1.32	2.02	2.24	3.56	1.77	2.52	2.72				
HEMFB1000490	9.25	6.82	8.08	12.41	16.92	19.33	9.89	8.92	10.33	*	+		
HEMFB1000491	6.31	3.37	4.57	9.52	7.65	10.48	5.02	4.69	3.79	*	+		
HEMFB1000492	2.22	0.64	1.44	4.93	5.13	7.41	2.99	2.91	3.63	**	+	*	+
HEMFB1000493	4.06	2.22	4.19	4.24	6.19	6.18	1.66	2.72	2.91				
HEMFB1000510	6.41	3.47	4.28	6.87	9.13	11.79	5.4	4.78	5.74				
HEMFB1000516	4.76	2.42	3.32	9.01	5.12	5.56	4.9	3.1	6.78				
HEMFB1000518	1.77	0.89	0.96	2.32	1.98	1.84	1.86	2.02	1.38				
HEMFB1000523	5.6	4.26	4.57	10.14	11.92	12.71	5.32	6.89	8.07	**	+		
HEMFB1000530	2.95	1.4	1.93	9.88	7.75	9.87	4.5	4.51	2.86	**	+		
HEMFB1000542	8.28	5.69	6.91	10.8	11.2	12.53	8.2	7.18	7.39	**	+		
HEMFB1000550	1.32	0.8	1.53	2.82	2.53	3.26	1.75	3.01	2.05	**	+		
HEMFB1000554	7.82	3.63	5.16	11.58	14.79	21.33	5.25	6.34	9.2	*	+		
HEMFB1000556	7.65	3.11	3.74	5.66	6.17	8.22	5.32	4.21	5.38				
HEMFB1000564	4.88	2.2	2.92	4.79	4.84	5.7	5.46	2.87	3.44				
HEMFB1000567	11.63	5.99	6.65	15.29	18.3	19.22	9.27	9.36	9.27	*	+		
HEMFB1000569	5.23	1.99	2.42	5.2	5.06	5.8	7.16	8.5	8.18		*	+	
HEMFB1000573	7.84	3.79	4.94	12.01	13.33	13.4	9.04	8.26	7.72	**	+		
HEMFB1000575	5.33	4.35	4.85	8.19	11.72	12.98	7.01	6.31	6.36	*	+	**	+

34/180

【0675】

【表204】

1545

1546

HEMBS1000579	1	0.63	1.23	1.94	1.94	1.61	0.75	2.27	0.83	*	+		
HEMBS1000583	1.32	0.9	1.33	2.89	2.66	2.35	1.39	2.41	1.82	**	+		
HEMBS1000586	5.03	2.33	2.86	4.93	10.49	10.9	3.19	3.33	3.66				
HEMBS1000589	4.34	3.31	2.32	4.73	9.62	7.86	4.05	4.47	4.07				
HEMBS1000591	6.2	2.47	3.35	5.53	10.43	9.55	5.26	4.88	5.68				
HEMBS1000592	3.62	1.12	1.49	3.68	3.48	4.83	5.06	2.83	3.4				
HEMBS1000593	5.63	3.16	4.14	7.95	8.98	9.6	4.23	4.57	4.71	**	+		
HEMBS1000594	9.73	4.88	6.49	11.51	8.83	10.26	5.12	4.65	3.54				
HEMBS1000598	3.08	2.45	2	3.88	5.18	4.28	2.89	4.3	2.68	*	+		
HEMBS1000611	1.33	0.64	1.43	2.46	1.17	1.82	0.83	1.24	1.6				
HEMBS1000617	12.12	5.56	4.61	11.59	16.06	19.06	7.94	6.34	10.83				
HEMBS1000633	7.8	2.76	2.97	7.01	3.89	6.02	4.57	2.94	6.61				
HEMBS1000630	2.59	1.28	1.39	2.17	2.39	2.78	2.69	2.13	3.79				
HEMBS1000631	10.27	4.76	4.53	6.2	6.77	8.48	8.04	7.46	8.07				
HEMBS1000632	6.25	2.1	3.02	6.63	6.59	8.13	4.84	4.67	4.51				
HEMBS1000636	13.35	4.72	8.11	7.29	10	13.28	8.71	9.3	9.58				
HEMBS1000637	26.51	17.46	16.75	28.37	43.24	52.91	24.53	21.76	22.76				
HEMBS1000638	1.76	0.67	1.19	2.95	4.12	4.45	1.31	0.92	1.68	**	+		
HEMBS1000642	10.59	4.41	5.99	11.15	12.92	13.73	6.73	6.84	9.2	*	+		
HEMBS1000643	1.65	1.83	1.24	2.38	2.51	3.19	2.28	0.92	1.97	*	+		
HEMBS1000649	3.91	2.47	2.78	5.9	5.23	6.96	3.56	3.95	5.15	*	+		
HEMBS1000652	6.02	2.91	2.8	5.46	7.5	7.04	3.21	3.43	4.33				
HEMBS1000655	12.28	6.34	8.07	9.28	11.26	11.56	6.56	3.92	6.25				
HEMBS1000665	1.52	0.76	1.22	2.5	1.48	1.81	2.25	0.85	1.56				
HEMBS1000668	2.21	0.39	1.35	5.91	7.44	6.43	4.09	4.69	4.22	**	+	**	+
HEMBS1000671	9.73	3.87	4.11	15	14.71	15.82	8.84	8.17	8.33	**	+		
HEMBS1000673	2	0.92	2.42	2.06	2.03	2.24	2.77	0.96	1.66				
HEMBS1000679	1.96	1.53	2.94	3.03	1.89	3.47	3.49	2.72	4.24				
HEMBS1000684	10.32	4.72	6.06	13.49	17.19	16.84	8.71	6	9.32	*	+		
HEMBS1000692	2.42	1.11	1.48	1.94	1.06	1.01	1.68	1.28	1.89				
HEMBS1000693	6.65	3.11	3.35	5.7	3.46	5.14	5.27	4.98	4.93				
HEMBS1000705	4.28	2.03	1.45	4.17	5.14	4.6	2.08	2.85	2.66				
HEMBS1000706	2.4	0.82	1.33	4.76	1.91	1.69	2.35	1.33	2.24				
HEMBS1000709	5.9	4.56	2.82	9.88	15.43	11.7	9.92	8.98	12.92	*	+	*	+
HEMBS1000714	4.07	1.84	2.28	3.51	2.48	3.46	4.34	1.56	2.5				
HEMBS1000725	3.83	2.12	2.8	3.51	3.57	2.91	4.38	2.17	3.5				
HEMBS1000726	6.74	3.26	3.37	8.38	10.66	11.11	5.09	6.26	5.9	*	+		
HEMBS1000729	5.92	3.12	3.67	3.82	5.2	5.28	2.93	3.03	3.74				
HEMBS1000738	6.27	2.98	4.84	7.01	7	9.14	5.8	4.68	8.01				
HEMBS1000749	6.38	4.5	8.03	10.82	12.38	19.82	6.87	7.43	9.13				
HEMBS1000763	4.28	1.52	4.69	3.87	3.73	4.04	3.58	5.24	3.54				
HEMBS1000770	2.56	1.54	1.45	4.69	5.02	5.12	3.94	2.82	2.01	**	+		
HEMBS1000774	4.01	2.16	2.61	6.02	5.76	6.03	4.48	3.56	3.59	**	+		
HEMBS1000777	16.82	8.94	10.71	11.64	9.96	10.04	11.16	9.95	10.48				
HEMBS1000781	4.68	2.51	2.03	4.83	6.62	5.74	2.82	4.66	5.27				
HEMBS1000788	1.26	1.09	0.22	0.77	1.4	0.96	0.82	1.05	1.38				
HEMBS1000789	3.3	1.16	1.77	2.42	1.9	2.76	1.89	2.74	1.95				
HEMBS1000790	4.72	2.05	3.39	5.79	6.37	7.78	3.19	2.91	4.28	*	+		
HEMBS1000794	0.97	0.54	1.08	1.04	2.04	2.15	0.72	1.24	1.02				
HEMBS1000807	7.3	3.23	3.76	7.53	4.81	6.34	3.19	2.77	3.98				
HEMBS1000809	10.2	3.24	6.13	7.78	12.54	11.13	7.52	8.8	9.69				
HEMBS1000810	6.83	2.64	2.68	4.19	3.73	4.74	4.18	2.82	5.16				
HEMBS1000821	3.04	1.01	1.43	1	1.91	2.05	1.27	2.15	1.75				
HEMBS1000822	1.16	1.15	0.89	1.14	1.34	1	1.68	1.67	1.68	**	+		
HEMBS1000826	3.27	2.25	2.9	2.37	8.91	8.1	2.85	5.14	2.76				
HEMBS1000827	4.04	1.85	2.66	4.07	6.2	5.58	3.55	3.41	2.85				

35/180

【0676】

【表205】

1548

HEMBA1000031	5.58	1.72	2.71	4.5	3.81	4.21	2.23	2.64	2.11								
HEMBA1000035	4	1.57	1.01	4.73	4.53	5.6	3.04	2.52	2.85	*	+						
HEMBA1000040	6.38	3.54	3.15	8.28	10.6	8.97	6.91	4.2	4.08	*	+						
HEMBA1000044	4.7	2.4	2.04	8.23	8.85	8.6	7.06	5.5	6.33	**	+	*	+	+	+	+	+
HEMBA1000052	0.54	0.28	0.27	0.52	0.36	0.24	1.16	0.97	0.61			*	+				
HEMBA1000057	7.91	6.39	3.23	5.68	6.47	7.09	4.42	3.6	4.37								
HEMBA1000058	5.33	2.35	2.78	9.3	8.37	8.17	3.94	3.82	2.97	**	+						
HEMBA1000067	5.01	2.6	3.3	9.23	10.12	8.69	3.49	5.17	4.43								
HEMBA1000070	4.43	1.73	2.81	6.64	6.44	7.5	2.8	3.34	3.99	*	+						
HEMBA1000076	2.52	1.01	1.78	2.03	2.41	3.32	1.17	1.96	2.6								
HEMBA1000081	4.52	2.25	2.68	3.85	3.48	4.21	3.8	3.6	3.52								
HEMBA1000083	1.07	0.87	0.48	2.38	2.52	2.42	1.86	2.24	1.15	**	+						
HEMBA1000087	16.17	10.38	8.54	18.39	28.8	26.71	14.31	15.73	15.23	*	+						
HEMBA1000089	1.52	0.47	0.72	0.71	0.87	1.25	1.08	2.34	2.95								
HEMBA1000090	4.2	1.91	2.82	6.2	6.22	11.04	3.56	3.57	3.05	*	+						
HEMBA1000093	3.13	1.95	2.57	3.14	8.44	5.73	3.88	3.35	2.73								
HEMBA1000098	2.72	1.85	1.78	2.31	2.75	4	1.77	1.83	1.88								
HEMBA1000095	7.13	4.79	4.05	6.15	5.33	7.36	6.49	7.74	6.04								
HEMBA1000098	3.42	1.78	2.53	3.45	3.15	4.99	2.18	3.31	2.95								
HEMBA1000099	3.27	1.5	0.99	3.5	4.25	4.18	2.64	2.6	2.61	*	+						
HEMBA10000913	1.53	1.02	1.16	2.35	1.71	3.01	2.43	2.82	3.12			**	+				
HEMBA10000915	125.5	96.58	90.74	52.7	70.12	78.2	138.4	94.57	151.2	*	-						
HEMBA10000917	5.94	3.71	3	10.02	9.8	10.14	6.41	5.43	5.2	**	+						
HEMBA10000927	3.9	2.3	4.04	2.93	2.18	2.45	3.26	2.61	3.09								
HEMBA10000933	1.41	0.52	1.78	2.08	2.21	2.86	1.55	1.9	0.46								
HEMBA10000933	63.34	47.44	31.38	44.11	52.4	49.52	46.54	37.21	45.55								
HEMBA10000936	7.16	3.79	4.04	4.95	3.87	5.38	3.06	2.19	2.36								
HEMBA10000939	9.8	5.4	5.5	8.13	8.11	6.88	7.11	4.16	5.78								
HEMBA10000941	1.26	1.52	1.91	2.33	1.33	3.43	1.03	2.28	3								
HEMBA10000947	3.84	2.12	3.17	3.27	3.95	6.16	2.65	3.42	5								
HEMBA10000954	2.09	0.96	1.77	3.22	2.47	2.01	1.52	2.5	2.09								
HEMBA10000959	1.47	0.69	1.99	4.15	4.21	5.2	2.08	3.64	2.15	**	+						
HEMBA10000973	0.93	0.22	1.08	1.36	1.53	1.02	0.58	1.34	0.88								
HEMBA10000975	6.35	2.45	2.52	2.87	4.55	4.7	3.97	3.56	3.46								
HEMBA10000981	1.55	0.65	1.17	2.92	1.74	2.12	1.91	1.15	1.6								
HEMBA10000985	4.16	2.16	3.38	6.79	6.53	7.43	6.9	5.56	5.46	**	+	*	+	+	+	+	+
HEMBA10000991	2.4	0.94	2.24	1.58	2.01	2.39	1.83	3.86	2.04								
HEMBA10000996	6.16	2.86	5.71	15.05	12.65	14.03	9.39	6.89	7.92	**	+						
HEMBA10010000	0.81	0.42	1.96	2.31	1.45	2	2.11	2.4	1.74								
HEMBA10010004	0.63	0.42	0.74	2.36	1.33	1.9	1.27	2.5	0.58	*	+						
HEMBA10010006	0.9	0.72	1.22	1.85	1.11	0.92	0.7	1.72	0.82								
HEMBA10010011	4.86	1.41	1.32	2.52	2.1	3.78	2.71	1.63	2.77								
HEMBA10020014	5.41	3.41	2.83	4.86	8.33	8.51	5.54	2.65	5.28								
HEMBA10010020	3.52	1.22	3.22	5.91	7.22	5.47	4.21	2.46	3.29	*	+						
HEMBA10010024	3.88	2.53	2.6	4.94	7.97	7.2	4.48	3.54	3.57	*	+						
HEMBA10010026	4.57	3.08	2.54	5.25	5.33	6.61	2.93	3.4	3.78								
HEMBA10010037	2.04	0.83	2.17	4.63	4.48	3.78	3.41	3.94	2.4	**	+						
HEMBA10010042	2.63	0.97	1.26	3.42	3.22	3.69	2.16	3.39	1.69	*	+						
HEMBA10010046	3.55	2.14	2.26	3.89	3.63	3.68	3.15	4.56	3.14								
HEMBA10010047	5	1.57	1.46	5.39	4.72	4.88	2.39	1.51	4.62								
HEMBA10010048	8.53	3.68	3.67	2.65	6.39	8.39	5.59	5.14	7.15								
HEMBA10010051	1.18	0.9	0.63	0.91	1.6	1.29	0.9	1.3	2.48								
HEMBA10010056	4.02	2.51	1.82	4.56	3.43	4.23	3.26	2.37	3.48								
HEMBA10010058	4.62	1.41	2.29	4.81	4.08	5.54	4.01	2.62	3.49								
HEMBA10010060	1.13	1.04	0.28	1.95	1.91	2.6	0.75	1.53	1.56	*	+						
HEMBA10010063	4.1	1.41	1.69	3.62	6.69	5.51	3.01	2.79	2.86								

1549

1550

HEMBS1001068	7.81	3.48	2.43	5.74	4.82	6.22	5.53	5.34	6.4										
HEMBS1001069	5.14	1.53	2.93	10.11	5.98	8.43	4.89	3.46	4.79	*									
HEMBS1001095	14.6	9.13	9.13	9.72	6.9	9.06	5.98	7.72	8.46										
HEMBS1001096	3.56	1.37	1.54	4.69	5.52	4.24	2.24	1.72	3.53	*									
HEMBS1001101	21.47	17.94	10.93	10.99	11.87	12.38	8.8	9.1	8.37										
HEMBS1001102	2.77	1.29	0.76	2.93	2.4	3.87	2.39	1.32	2.26										
HEMBS1001104	5.43	2.94	3.94	9.11	5.73	9.85	5.68	2.83	4.42	*									
HEMBS1001105	3.73	2.54	3.47	3.95	6.18	9.09	3.99	3.81	3.94										
HEMBS1001112	8.37	6.64	4.97	5.94	6.55	6.82	6.29	6.97	5.99										
HEMBS1001113	7.58	3.55	4.62	10.53	11.56	12	7.39	5.47	7.82	**									
HEMBS1001114	7.84	3.54	5.33	11.15	12.39	11.97	6.57	3.79	5.53	**									
HEMBS1001115	12.69	6.52	6.38	8.41	6.32	7.74	8.1	3.98	5.13										
HEMBS1001117	1.26	0.59	1.14	3.99	3.99	7.09	4.39	3.19	2.89	*									
HEMBS1001119	2.73	0.69	1.36	3.27	2.76	3.17	1.69	1.82	2.33										
HEMBS1001126	17.3	8.41	6.34	12.51	13.52	16.39	9.04	9.96	8.26										
HEMBS1001133	7.22	2.46	6.43	7.94	11.25	15.48	5.58	5.96	7.46										
HEMBS1001137	4.69	1.94	2.48	3.07	2.31	3.24	4.3	2.74	3.49										
HEMBS1001142	10.97	4.26	5.7	14.69	16.82	16.36	7.91	5.78	10.87	*									
HEMBS1001145	8.34	3.24	4.81	10.74	10.95	12.08	5.82	4.69	6.65	*									
HEMBS1001151	8.95	6.02	5.47	5.12	6.22	5.78	8.43	8.19	8.82										
HEMBS1001153	5.68	3.55	3.85	6.9	7.36	7.26	5.29	4.07	4.12	*									
HEMBS1001158	5.25	4.46	4.73	8.21	9.2	10.97	4.6	4.37	5.83	**									
HEMBS1001162	5.93	2.46	2.66	6.12	6.91	7.13	3.71	3.73	4.71										
HEMBS1001170	2.28	0.23	1.68	2.09	1.33	2.33	1.48	1.17	1.14										
HEMBS1001175	4.7	2.5	2.14	5.28	3.05	6.25	4.06	3.09	3.56										
HEMBS1001177	11.32	4.92	7.58	14.33	14.36	15.14	8.51	7.62	8	*									
HEMBS1001182	7.1	3.3	3.03	8.51	7.41	6.84	6.75	4.9	5.74										
HEMBS1001192	4.01	1.43	2.59	3.22	2.9	2.65	3.81	3.22	2.43										
HEMBS1001199	1.24	0.85	1.37	0.51	1.77	3.72	1.58	1.98	1.27										
HEMBS1001200	0.7	0.28	0.37	0.41	0.29	1.06	0.14	0.69	0.72										
HEMBS1001208	6.24	1.58	2.41	2.54	3.62	5	2.67	3.31	3.15										
HEMBS1001209	8.96	2.6	4.27	8.47	9.46	10.64	6.12	3.72	4.78										
HEMBS1001210	3.39	3.6	6.25	13.57	15.06	13.24	8.2	7.86	10.28	**									
HEMBS1001215	56.1	31.37	29.04	36.73	42.52	41.17	25.87	19.36	26.75										
HEMBS1001217	4.33	2.5	3.14	2.96	3.91	4.21	4.42	3.57	4.01										
HEMBS1001218	4.39	2.08	2.28	6.07	7.97	8.92	4.93	4.87	4.51	*									
HEMBS1001221	1.61	1.15	0.66	1.21	1.16	1.19	2.11	1.68	0.87										
HEMBS1001224	2.88	1.37	1.83	3.46	3.87	4.78	1.63	2.85	1.71	*									
HEMBS1001230	3.6	1.44	3.39	4.28	5.22	5.68	2.22	3.15	2.2	*									
HEMBS1001234	9.13	2.44	8.29	5.98	6.49	5.96	5.83	7.02	6.04										
HEMBS1001235	5.3	2.57	3.09	3.97	3.82	5.68	4.42	5.01	5.46										
HEMBS1001237	11.86	5.88	6.73	9.88	9.37	10.19	7.04	5.53	6.3										
HEMBS1001242	3.75	2.48	2.08	4.97	4.37	4.59	4.13	4.47	3.96	*									
HEMBS1001244	1.32	1.13	0.4	0.82	0.94	1.53	1.73	1.61	1.2										
HEMBS1001249	3.12	1.54	0.34	2.25	4.83	2.55	2.05	1.99	2.11										
HEMBS1001253	6.29	1.42	2.97	13.67	4.84	8.24	2.79	2.84	4.65										
HEMBS1001254	2.47	0.84	1.05	1.37	2.56	1.79	1.57	2.54	1.52										
HEMBS1001266	1.23	0.44	1.59	2.72	2.03	1.62	2.12	1.69	5.16										
HEMBS1001267	7.87	4.02	4.63	13.5	11.84	13.24	5.84	7.38	7.83	**									
HEMBS1001271	4.61	1.62	1.38	4.06	3.96	5.87	2.53	2.67	2.49										
HEMBS1001282	6.27	3.11	3.61	3.44	3.72	3.72	3.96	3.68	3.39										
HEMBS1001287	13.66	7.12	7.62	9.05	8.08	10.38	11.92	6.12	11.75										
HEMBS1001288	3.65	1.71	2.11	2.38	2.54	2.69	2.09	2.39	1.58										
HEMBS1001289	10.93	6.03	8.57	15.81	16.55	18.1	8.4	7.7	5.92	**									
HEMBS1001290	3.6	2.6	2.26	3.2	3.1	5.19	4.49	3.29	2.53										
HEMBS1001294	2.74	1.82	3.02	1.97	1.99	2.92	2.55	2.49	2.42										

37/180

【0678】

【表207】

1551

1552

HEMBS1001299	11.58	8.15	6.05	9.03	8.73	7.82	6.87	6.29	8.87										
HEMBS1001302	6.82	4.33	3.28	5.31	5.44	7.1	4.47	4.4	7.2										
HEMBS1001304	1.87	0.87	0.83	1.3	1.76	2.94	1.91	2.12	1.37										
HEMBS1001314	2.52	0.38	1.35	1.89	2.07	2.7	1	1.94	1.9										
HEMBS1001315	2.2	0.42	0.99	1.6	0.99	1.5	2.82	1.66	1.14										
HEMBS1001317	5.5	2.93	3.71	6	6.29	9.01	6.12	6.25	7.04										
HEMBS1001326	1.44	0.28	0.42	1.14	1.37	0.97	0.93	1.85	0.56										
HEMBS1001331	3.49	1.15	3.33	3.16	5.21	4.92	2.94	2.32	3.4										
HEMBS1001335	2.13	0.58	1.32	2.09	1.33	1.73	1.47	0.84	0.69										
HEMBS1001337	4.69	2.11	3.26	4.29	6.51	6.35	3.43	4.14	3.13										
HEMBS1001339	3.42	1.11	1.36	2.82	1.69	2.07	1.52	2.17	1.96										
HEMBS1001344	2.99	1.77	1.84	2.4	2.3	3.56	2.28	2.27	2.05										
HEMBS1001346	3.15	2.58	2.53	3.75	3.57	4.79	2.76	4.39	3.22	*									
HEMBS1001348	1.96	1.25	1.97	4.3	3.56	4.57	1.75	3.51	2.64	**									
HEMBS1001350	2.69	1.8	2.82	11.17	12.63	10.95	7.44	8.11	8.86	**									
HEMBS1001356	1.82	0.34	1.21	2	1.23	1.35	0.99	1.53	1.63										
HEMBS1001364	1.29	0.93	0.89	1.8	2.27	2.41	2.29	1.24	1.25	**									
HEMBS1001366	3.41	1.36	1.76	6.29	5.97	7.89	2.97	3.23	3.76	**									
HEMBS1001367	5.44	2.63	4.67	5.82	13.11	9.17	6.34	5.62	5.1										
HEMBS1001369	1.88	0.36	0.91	2.5	3.44	2.87	2.19	3.7	2.34	*									
HEMBS1001380	3.65	2.5	3.07	8.69	9.13	10.12	4.6	7.63	4.24	**									
HEMBS1001381	7.54	3.35	4.95	9.78	7.21	8.91	5.88	6.12	6.67										
HEMBS1001384	2.77	2.23	5.27	4.04	4.7	5.21	2.99	3.46	4										
HEMBS1001387	1.33	0.72	1.19	2.84	1.92	3.26	0.78	2.08	0.69	*									
HEMBS1001394	2.01	1.22	0.71	4.71	4.19	4.99	2.39	2.44	2.66	**									
HEMBS1001407	3.37	1.49	0.8	2.53	3.21	2.87	4.47	1.2	2										
HEMBS1001410	1.19	0.14	0.37	0.55	0.79	0.77	0.44	1.14	0.17										
HEMBS1001413	2.53	1.15	2.11	4.01	6.2	3.82	2.17	2.18	2.56	*									
HEMBS1001419	3.82	1.67	2	5.53	5.54	4.76	5.16	3.44	3.45	*									
HEMBS1001421	1.55	0.78	1.24	9.94	7.28	9.56	5.74	5.75	4.91	**									
HEMBS1001424	0.54	0	0.28	0.9	0.45	0.6	0.84	1.22	0.47										
HEMBS1001426	2.45	0.64	1.42	3.9	4.18	3.95	2.09	3.09	1.9	**									
HEMBS1001429	10.12	5.99	4.62	6.28	4.44	8.1	5.21	7.29	9.1										
HEMBS1001436	11.8	4.02	6.29	22.88	14.63	21.79	9.57	8.07	10.97	*									
HEMBS1001443	1.46	1.5	1.3	2.55	2.11	3.84	5.74	4.67	5.74										
HEMBS1001449	4.24	1.68	1.33	4.21	5.76	5.46	2.38	1.89	2.76										
HEMBS1001454	4.2	2.22	2.85	4.88	5.14	6.3	1.94	2.02	3.61	*									
HEMBS1001458	4.34	4.36	3.05	7.92	4.69	4.55	3.87	3.06	3.94										
HEMBS1001461	2.41	1.63	1.39	3.76	3.78	6.76	3.87	1.93	2.34	*									
HEMBS1001463	4.41	1.84	3.33	6.77	8.03	7.56	3.07	2.66	3.3	**									
HEMBS1001464	1.53	1.48	0.96	1.16	0.81	1	0.81	0.25	1.04										
HEMBS1001466	1.71	1.2	0.87	3.03	2.72	4.34	2.85	2.09	4.25	*									
HEMBS1001482	3.03	1.42	1.06	1.64	2.18	1.42	2.97	1.16	2.1										
HEMBS1001500	2.17	1.05	0.9	2.57	2.02	2.37	1.04	1.45	1.55										
HEMBS1001505	8.22	5.06	7.49	13.32	13.9	13.27	5.5	6.16	7.01	**									
HEMBS1001521	2.58	1.03	1.95	4.68	3.52	3.79	2.8	2.46	2.3	*									
HEMBS1001527	14.66	7.32	7.32	12.93	16.36	15.19	7.53	11.09	12.62										
HEMBS1001530	7.24	3.1	6.46	5.19	6.93	5.94	6.69	5.92	5.53										
HEMBS1001531	5.66	2.3	2.38	5.05	4.74	5.69	3.58	2.66	2.99										
HEMBS1001532	2.05	0.38	0.82	1.99	0.87	2.3	1.76	1.25	1.24										
HEMBS1001535	3.86	2.42	2.26	4.62	4.93	5.74	3.17	2.1	4.36	*									
HEMBS1001536	5.02	2.43	2.77	5.57	4.42	5.08	2.95	2.46	3.39										
HEMBS1001537	3.43	1.79	1.93	5.9	3.91	6.35	3.35	2.86	3.81	*									
HEMBS1001542	10.24	4.77	6.29	8.68	10.49	11.37	4.75	4.74	4.61										
HEMBS1001543	4.42	2	4.45	6.17	7.07	7.41	4.96	3.35	2.51	*									
HEMBS1001547	1.69	0.68	1.1	3.41	2.74	1.36	1.07	2.16	2.08										

38/180

【0679】

【表208】

1553

1554

HEMBS1001548	11.61	4.55	5.07	6.22	6.23	8.02	13.1	5.3	6.57										
HEMBS1001549	2.02	1.27	1.35	2.89	1.88	2.65	2.33	1.18	2										
HEMBS1001550	3.38	2.36	2.27	4.34	5.15	4.75	3.71	2.88	3.52	**	+								
HEMBS1001551	6.73	3.72	2.72	6.03	4.98	4.88	4.48	4.21	3.29										
HEMBS1001552	143.7	103.7	84.35	117.1	130.5	149.5	73.73	78.89	81.45										
HEMBS1001553	4.34	2.01	6.14	5.35	4.71	6.93	2.17	3.81	2.88										
HEMBS1001554	3.35	1.85	2.92	2.44	1.48	3	2.68	1.66	2.47										
HEMBS1001555	4.11	1.78	1.25	2.55	2.8	4.22	1.76	2.85	2.7										
HEMBS1001556	5.19	3.43	2.13	7.14	9.58	10.48	5.06	4.35	4.95	*	+								
HEMBS1001557	2.45	1.89	1.47	2.45	2.59	4.79	2.08	2.01	1.8										
HEMBS1001558	9.91	4.02	1.68	7.84	12.44	10.86	6.94	6.15	5.93										
HEMBS1001559	2.38	2.13	1.24	3.04	4.7	3.31	4.54	3.91	4.77	*	**	+							
HEMBS1001560	7.58	3.68	4.12	10.26	11.71	11.73	8.26	5.8	7.17	**	+								
HEMBS1001561	1.66	1.47	1.01	2.08	1.72	2.54	1.43	2.23	1.83										
HEMBS1001562	1.5	0.25	0.77	1.78	2.38	2.95	1.47	2.06	1.36	*	+								
HEMBS1001563	0.98	0.3	0.79	0.72	0.7	0.98	0.73	0.96	0.76										
HEMBS1001564	7.29	5.01	5.69	10.05	12.84	11.6	6.84	5.75	5.35	**	+								
HEMBS1001565	2.21	1.9	1	2.28	2.95	2.82	2.58	3.52	1.79										
HEMBS1001566	2.74	2.34	1.59	5	7.12	6.26	2.86	3.86	3.26	**	+								
HEMBS1001567	3.47	2.37	1.26	9.12	1.21	1.26	2.81	2.15	1.28										
HEMBS1001568	0.39	0.5	0.61	1.56	1.46	2.32	2.13	1.91	2.02	*	**	+							
HEMBS1001569	2.05	0.69	1.57	1.73	2.03	1.92	0.69	0.97	1.11										
HEMBS1001570	2.2	0.75	1.17	3.5	2.23	1.77	1.56	1.05	1.51										
HEMBS1001571	3.51	1.4	2.57	3.58	4.43	4.86	2.1	2.95	2.6										
HEMBS1001572	1.95	0.54	0.63	1.54	1.04	1.19	1.35	0.64	1.26										
HEMBS1001573	5.49	2.4	2.56	5.29	5.68	6.05	3.35	3.68	4.27										
HEMBS1001574	1.36	1.13	0.8	0.24	0.85	0.87	0.48	0.61	0.56	*	-								
HEMBS1001575	2.05	1.95	2.11	3.16	2.96	2.94	3.08	3.78	1.71	**	+								
HEMBS1001576	2.49	2.15	1.55	5.36	1.62	4.96	1.46	1.39	2.66										
HEMBS1001577	1.24	0.08	2.02	7.77	6.22	7.71	3.16	4.06	4.45	**	+	*	+						
HEMBS1001578	1.14	0.56	0.64	1.01	1.36	1.96	0.82	0.73	1.12										
HEMBS1001579	4.9	1.43	3.88	3.76	6.22	5.35	4.26	5.92	6.05										
HEMBS1001580	9.43	4.46	3.65	7.18	5.87	10.36	4.73	4.98	5.54										
HEMBS1001581	4.45	1.52	2.55	2.96	2.17	2.25	2.39	2.98	2.34										
HEMBS1001582	3.43	1.92	1.36	3.15	2.26	1.5	2.37	3.04	2.3										
HEMBS1001583	3.34	2.15	1.93	2.33	2.97	3.86	3.7	4.19	3.07										
HEMBS1001584	0.43	0.79	0.82	2.14	2.22	2.08	1.31	1.84	3.11	**	+								
HEMBS1001585	0.91	0	0.49	2.21	2.23	2.38	1.38	2.34	1.74	**	+	*	+						
HEMBS1001586	8.08	2.9	6.21	6.72	7.83	9.08	5.46	5.54	6.16										
HEMBS1001587	4.34	1.92	3.68	6.91	10.28	8.29	3.81	4.11	3.01	*	+								
HEMBS1001588	5.33	4.91	1.92	7.82	8.35	10.07	3.64	3.58	4.65	*	+								
HEMBS1001589	5.79	3.89	4.11	6.65	6.58	7.83	3.81	4.55	4.31	*	+								
HEMBS1001590	2.9	1.19	1.54	2.56	2.78	3.16	1.54	2.75	1.87										
HEMBS1001591	36.41	33.52	31.32	28.11	26.74	25.14	22.32	14.43	19.67	*	-	**	-						
HEMBS1001592	3.1	2.92	2.47	4.7	5.73	6.77	4.4	3.15	4.26	**	+								
HEMBS1001593	2.54	0.66	2.3	5.4	4.73	5.13	2.48	3.07	2.77	**	+								
HEMBS1001594	5.75	4.06	4.43	6.69	5.77	10.19	4.22	3.36	4.39										
HEMBS1001595	2.44	0.77	1.23	3.44	4.21	3.48	1.4	1.37	2.22	*	+								
HEMBS1001596	8.77	3.39	4.72	11.21	15.68	17.47	7.58	5.43	7.33	*	+								
HEMBS1001597	7.34	3.22	3.36	7.29	7.53	8.22	6.29	5.11	5.25										
HEMBS1001598	3.12	1.84	2.45	2.82	2.94	4.26	3.19	1.55	3.2										
HEMBS1001599	0.84	0.19	0.52	0.79	1.64	1.25	0.88	2.16	1.24										
HEMBS1001600	1.15	0.71	0.71	1.53	1.28	1.9	0.58	2.3	0.49	*	+								
HEMBS1001601	2.92	1.03	2.15	2.66	1.94	2.8	2.3	3.2	2.05										
HEMBS1001602	11.82	11.49	14.29	11.74	11.14	12.3	6.31	7.75	11.3										
HEMBS1001603	0.42	0.01	1.19	1.62	1.09	1.43	0.08	1.04	1.6										

39/180

【0680】

【表209】

1555

1556

HEMBS1001788	5.11	2.85	2.49	8.04	8.23	9.77	5.27	5.14	5.83	**	+		
HEMBS1001789	13.59	3.52	4.92	5.61	7.12	5.14	6.71	5.28	5.84				
HEMBS1001797	0.88	0.62	1.95	0.94	0.65	0.97	1.07	1.81	1.9				
HEMBS1001802	6.5	3.72	4.06	7.4	8.03	6.58	5.93	6.91	6.06				
HEMBS1001812	5.74	3.61	5.29	9.39	12.73	12.64	5.58	6.99	9.37	**	+		
HEMBS1001815	20.05	9	15.52	27.98	23.86	26.02	37.42	29.06	44.83	*	+	*	+
HEMBS1001816	5.07	2.26	3.92	9.09	8.62	9.45	5.29	4.77	4.9	**	+		
HEMBS1001831	1.2	0.45	0.53	1.8	1.74	1.99	0.55	2.73	1.28	*	+		
HEMBS1001834	19.83	12.47	10.64	12.5	19.26	19.83	14.74	13.9	15.71				
HEMBS1001836	4.06	3.15	2.68	7.01	7.21	7.9	3.1	3.18	4.01	**	+		
HEMBS1001839	1.83	0.36	0.78	1.33	1.05	1.21	1.58	1.39	1.02				
HEMBS1001841	4.21	3.05	4.61	6.62	7.34	6.85	8.41	7.68	5.57	**	+	*	+
HEMBS1001844	4.31	2.59	2.19	5.78	3.8	4.04	2.62	4.06	3				
HEMBS1001847	11.75	7.18	10.2	21.65	17.41	24.55	7.68	9.92	9.98	**	+		
HEMBS1001848	2.73	1.25	1.47	4.72	2.91	3.06	15.56	19.7	16.79		**	+	
HEMBS1001850	7.3	4.6	5.92	9.74	8.83	8.43	10.59	7.86	13.13	*	+		
HEMBS1001852	6.4	9.16	9.93	12.13	14.98	16.02	18.07	14.33	23.47	*	+	*	+
HEMBS1001863	6.66	2.82	3.58	9.9	10.12	11.35	6.68	3.16	7.05	**	+		
HEMBS1001867	1.21	1.36	0.82	2.34	2.45	3.53	2.08	1.31	1.98	*	+		
HEMBS1001868	3.28	1.27	0.26	2.34	1.83	1.98	2.3	1.36	2				
HEMBS1001869	4.99	3.41	2.47	4.55	8.08	7.57	3.34	3.94	4.05				
HEMBS1001872	3.4	4.06	0.84	4.75	2.37	1.57	2.65	1.38	2.04				
HEMBS1001874	2.47	1.57	1.58	3.42	1.79	3.58	3.5	1.76	2.08				
HEMBS1001875	1.3	0.4	3.1	2.27	2.57	2.84	2.23	0.73	0.98				
HEMBS1001880	9.6	4.1	4.24	11.57	10.59	10.4	5.78	4.19	6.8				
HEMBS1001889	2.12	0.58	0.29	1.53	1.49	1.79	2.01	0.55	1.92				
HEMBS1001903	4.86	1.84	3.46	4.45	3.55	4.47	5.08	3.38	4.99				
HEMBS1001905	6.94	3.72	4.24	3.83	3.28	4.45	3.35	1.95	3.04				
HEMBS1001906	3.51	0.89	1.09	3.56	2.45	3.39	2.27	3.05	2.44				
HEMBS1001908	1.61	2.17	1.92	5.17	4.2	3.43	1.41	2.29	3.44	*	+		
HEMBS1001910	2.88	1.38	0.82	4.07	3.93	6.71	2.4	1.88	3.55	*	+		
HEMBS1001911	6.98	2.87	4.02	9.07	10.54	12.95	3.98	4.78	7.22	*	+		
HEMBS1001915	4.25	1.76	1.83	6.42	5.24	7.19	5.74	2.92	4.49	*	+		
HEMBS1001921	5.38	3.56	4.5	10.21	11.3	11	5.97	4.64	6.62	**	+		
HEMBS1001922	3.83	1.35	3.8	5.95	3.77	3.39	3.48	2.3	3.67				
HEMBS1001923	3.73	2.29	2.11	4.2	3.69	3.62	2.81	2.27	3.72				
HEMBS1001930	0.59	0.63	0.42	2.23	1.25	1.36	0.41	1.35	1.01	*	+		
HEMBS1001944	3.88	3.55	3.94	5.26	8.37	10.06	2.98	4.95	4.6	*	+		
HEMBS1001945	5.17	3.58	5.47	3.15	4.34	6.51	3.41	6.48	6.46				
HEMBS1001947	6.49	1.48	5.58	2.11	3.59	4.92	2.72	2.7	2.62				
HEMBS1001950	6.47	3.08	4.75	4.98	5.8	5.65	5.08	4.12	4.55				
HEMBS1001952	4.62	1.75	2.38	5.87	7.63	6.22	3.88	3.07	2.9	*	+		
HEMBS1001953	3.33	1.23	1.69	3.8	4.29	3.6	2.72	2.28	2.79				
HEMBS1001957	3.22	1.56	1.85	3.38	4.52	4.53	3.81	1.96	3.18	*	+		
HEMBS1001959	7.02	7.17	6.24	7.94	4.73	8.54	5.15	5.79	4.06		*	-	
HEMBS1001963	4.04	1.76	3.14	4.32	4.25	6.26	2	2.46	3.87				
HEMBS1001967	11.44	5.2	6.57	12.83	15.13	16.73	7.11	8.82	7.6	*	+		
HEMBS1001973	5.08	2.32	4.1	4.86	6.84	9.36	3.18	5.36	3.55				
HEMBS1001978	7.53	3.35	6.01	7.28	6.5	6.97	5.53	5.59	5.54				
HEMBS1001983	20.88	11.32	14	10.33	15.1	15.82	11.27	9.2	12.69				
HEMBS1001987	1.67	0.99	0.76	2.21	2.1	2.79	2.2	1.61	1.87	*	+		
HEMBS1001988	1.86	1.73	2.04	2.83	3.58	3.43	2.01	2.02	1.85	**	+		
HEMBS1001990	4.63	2.51	4.22	4.26	6.45	6.12	4.77	3.77	5.16				
HEMBS1001996	2.64	1.19	1.29	1.17	1.43	2.67	1.72	2.23	1.63				
HEMBS1001997	4.3	2.22	2.71	5.89	6.32	7.41	2.43	4.16	2.74	*	+		
HEMBS1001999	15.97	11.41	12.12	8.02	17.07	19.1	5.81	7.78	7.71		*	-	

40/180

【0681】

【表210】

1557

1558

HEMBB1002002	0.83	0.59	1.4	1.42	1.71	2.28	1.59	0.62	1.07										
HEMBB1002003	8.43	2.74	4.65	11.77	11.48	12.25	6.14	5.42	7.41	*	+								
HEMBB1002009	0.77	2.18	1.38	1.23	1.38	2.16	0.85	1.5	0.79										
HEMBB1002013	2.33	1.35	1.79	1.62	1.81	3.45	1.55	1.11	1.73										
HEMBB1002015	7.48	4.38	3.67	9.87	8.21	13.87	7.92	7.55	9.97										
HEMBB1002024	12.18	6.96	6.46	7.22	8.12	8.32	6.32	9.38	7.79										
HEMBB1002033	3.12	1.84	1.81	4.86	5.45	3.22	1.97	2.8	1.44	*	+								
HEMBB1002039	3.05	1.27	3	3.79	6.93	5.61	2.49	3.6	2.96										
HEMBB1002041	7.09	2.89	3.99	5.42	7.13	7.97	5.81	4.83	6.2										
HEMBB1002042	7.43	3.78	4.66	7.93	11.47	10.08	5.53	6.71	6.67	*	+								
HEMBB1002043	4.31	1.3	3	5.84	8.07	8.67	4.27	3.64	4.54	*	+								
HEMBB1002044	1.54	1.29	1.16	1.41	1.89	1.39	1.48	2.04	1.41										
HEMBB1002045	13.56	9.28	9.85	18.7	19.69	19.62	11.33	11.49	14.07	**	+								
HEMBB1002049	0.94	0.9	1.48	2.03	3.05	3.51	1.86	1.85	1.5	*	+	*	+						
HEMBB1002050	2.63	0.87	2.41	2.24	3.31	3.77	1.82	2.42	2.29										
HEMBB1002051	2.77	1.42	2.72	3.76	4.08	3.57	1	2.97	1.66	*	+								
HEMBB1002058	11.05	4.29	3.63	7.71	6.55	7.57	7.7	4.29	6.63										
HEMBB1002069	13.1	6.94	8.01	16.77	20.06	18.1	11.13	9.92	13.2	*	+								
HEMBB1002075	2.31	1.12	2.72	4.01	5.39	4.96	2.61	2.52	2.47	*	+								
HEMBB1002079	3.29	1.28	2.08	2.22	2.42	2.23	2.53	2.39	1.66										
HEMBB1002080	1.83	2.55	0.96	2.15	2.98	4.39	1.68	2.81	2.3										
HEMBB1002082	2.22	1.44	1.38	1.35	2.4	2.6	1.2	1.53	2.07										
HEMBB1002084	1.85	1.72	1.79	2.73	3.83	5.21	2.72	3.71	3.91	*	+	*	+						
HEMBB1002088	11.64	8.26	10.3	14.66	19.71	16.32	16.11	15.05	19.56	*	+	*	+						
HEMBB1002092	8.42	4.12	3.19	8.1	10.6	9.29	6.67	5.28	5.88										
HEMBB1002094	8.51	6.18	7.26	14.48	15.44	15.77	7.48	6.89	8.09	**	+								
HEMBB1002103	13.1	13.5	12.83	61.49	68.55	57.48	66.63	34.04	51.6	**	+	*	+						
HEMBB1002109	6.77	3.65	4.41	10.27	12.78	11.5	7.97	4.24	7.06	**	+								
HEMBB1002115	44.63	28.15	32.39	41.8	53.57	63.47	24.84	22.28	27.42										
HEMBB1002120	2.22	0.77	1.3	3.55	2.83	2.5	1.74	2.54	1.48	*	+								
HEMBB1002121	1.32	0.72	1.59	2.14	1.84	1.52	1.15	1.56	1.25										
HEMBB1002134	29.98	14.03	18.39	22.56	28.18	29.08	20.1	20.18	26.29										
HEMBB1002136	5.67	2.48	3.78	3.62	3.43	4.97	3.89	4.13	4.88										
HEMBB1002138	3.55	2.31	2.47	7.41	6.73	5.61	7.6	5.28	8.06	**	+	*	+						
HEMBB1002139	3.56	2.49	3.1	6.05	5.07	6.19	3.34	5.1	3.14	**	+								
HEMBB1002141	5.57	2.73	5.33	5.02	6.05	7.64	4.99	5.45	6.15										
HEMBB1002142	4.26	2.17	2.9	5.21	4.83	7.21	3.06	3.4	2.29										
HEMBB1002145	2.66	1.68	2.79	4.87	2.84	2.91	1.83	3.33	2.18										
HEMBB1002152	2.89	1.29	3.31	6.08	5.5	7.8	2.66	3.88	3.38	*	+								
HEMBB1002162	4.47	2.09	2.74	4.63	5.63	4.42	2.84	4.52	4.28										
HEMBB1002173	2.01	1.5	1.53	4.12	5.2	7.12	2.21	2.47	3.85	*	+								
HEMBB1002189	5.63	4.01	3.4	9.38	12.87	12.35	5.18	5.01	5.41	**	+								
HEMBB1002190	4.01	6.72	3.24	8.35	6.45	9.57	5.06	3.62	5.05										
HEMBB1002193	4.3	2.37	3.54	3.79	4.24	4.57	3.11	3.84	2.85										
HEMBB1002217	8.31	4.18	4.51	10.88	11.96	11.17	4.63	6.35	5.39	*	+								
HEMBB1002218	21.17	7.63	13.71	19.12	24.55	22.92	14.78	13.7	19.7										
HEMBB1002228	4.29	2.39	3.53	7.69	9.04	7.22	3.92	7.05	4.8	**	+								
HEMBB1002232	2.54	0.96	2.12	5.44	4.77	4.87	2	4.33	3.46	**	+								
HEMBB1002245	2.24	0.69	1.25	1.7	1.97	1.97	1.7	1.11	1.6										
HEMBB1002247	2.78	1.52	2.56	1.84	2.86	2.26	3.27	2.52	2.93										
HEMBB1002249	8.45	3.73	4.77	12.48	12.32	13.64	6.18	5.35	5.42	**	+								
HEMBB1002254	2.12	1.02	1.52	4.72	4.67	7.07	3.96	3.27	2.9	**	+	*	+						
HEMBB1002255	0.31	0.16	1.07	0.59	0.84	2.46	0.5	0.92	0.27										
HEMBB1002266	1.03	0.51	0.66	4.13	2.54	2.5	1.25	1.72	1.36	*	+	*	+						
HEMBB1002271	56.56	35.65	38.07	20.53	29.91	28.83	14.93	16.36	14.09	*	-								
HEMBB1002280	1.89	0.47	1.28	2.71	3.38	2.75	1.12	1.95	1.11	*	+								

41/180

【0682】

【表211】

1559

1560

HEMFB1002296	19.39	12.59	10.1	13.85	9.77	11.58	17.38	19.09	20.6										
HEMFB1002300	5.98	2.27	2.27	4.97	4.83	5.06	3.39	2.79	3.87										
HEMFB1002302	4.79	2.37	2.24	3.34	4.96	4.22	3.13	3.11	2.5										
HEMFB1002306	2.53	0.59	1.19	2.93	4.01	3.53	2.16	2.15	1.9	*									
HEMFB1002316	1.37	0.21	1.01	1.03	1.85	1.65	1.5	1.08	0.63										
HEMFB1002326	9.34	4.41	4.08	6.83	11.8	13.52	5.14	6.95	5.58										
HEMFB1002327	3.74	1.52	2.2	3.25	6.69	8.05	1.41	2.57	2.14										
HEMFB1002339	6.65	2.85	3.03	3.55	3.52	4.81	3.39	4	4.24										
HEMFB1002340	2.45	1.14	0.8	2.72	7.22	1.38	1.47	2.32	1.56										
HEMFB1002342	18.78	10.67	11.1	11.48	10.39	11.81	11.37	10.6	12.37										
HEMFB1002348	8.06	4.65	5.88	8.32	11.43	13.39	7.12	5.37	8.32										
HEMFB1002349	4.65	2.7	3.21	2.57	3.59	5.52	2.08	3.08	3.75										
HEMFB1002364	3.68	2.01	1.94	4.35	5.19	5.2	2.34	2.77	3.18	*									
HEMFB1002366	26.64	15.48	15.83	13.61	16.98	21.16	15.49	15.51	17.68										
HEMFB1002371	2.23	1.84	1.61	9.83	11.88	12.25	6.86	8.63	9.93	**		+	**						
HEMFB1002381	6.41	3.55	2.93	4.03	6.29	6.16	5.19	4.39	5.77										
HEMFB1002383	10.2	4.93	4.09	9.89	9.52	10.26	9.31	9.32	10.54										
HEMFB1002387	11.72	4.82	7.2	7.69	8.97	9.71	6.05	7.95	7.6										
HEMFB1002389	4.35	2.96	2.55	5.95	6.17	9.26	3.8	3.76	3.86	*									
HEMFB1002413	10.96	4.94	5.84	12.47	15.22	15.46	7.04	7.35	7.5	*									
HEMFB1002415	2.9	1.63	1.04	2.46	1.99	2.7	2.07	2.58	1.35										
HEMFB1002424	2.41	2.37	3.44	2.94	2.65	3.7	0.8	2.25	2.17										
HEMFB1002435	6.05	3.85	3.42	8.18	9.21	12.24	4.22	6.67	5.02	*									
HEMFB1002437	8.18	4.1	4.67	3.14	4.27	5.26	6.03	4.48	3.96										
HEMFB1002443	12.17	4.35	6.23	11.86	16.23	14.17	10.19	3.68	8.32										
HEMFB1002447	8.82	3.51	5.23	10.28	11.65	12.71	5.54	5.46	6.69	*									
HEMFB1002453	10.1	3.7	4.44	12.22	12.96	16.08	5.85	7.3	7.02	*									
HEMFB1002457	8.34	2.86	3.7	8.87	9.3	9.53	4.63	5.01	4.51										
HEMFB1002458	1.84	0.2	0.83	2.21	1.65	2.32	1.18	4.23	1.99										
HEMFB1002463	13.99	7.17	7.29	17.97	18.05	72.29	8.48	10.09	10.66	*									
HEMFB1002465	3.55	1.09	2.46	1.87	2.68	3.41	1.36	3	1.53										
HEMFB1002477	3.8	1.74	1.62	2.44	2.7	2.39	2.93	1.14	1.8										
HEMFB1002479	1.35	1.53	2.03	10.77	11.28	12.82	19.91	17.51	11.35	**		+	**						
HEMFB1002482	8.63	4.67	6.83	7.48	7.18	7.8	5.28	6.57	5.43										
HEMFB1002492	2.72	1.93	0.73	4.53	3.38	4.56	3.26	3.14	4.65	**		+	**						
HEMFB1002495	5.34	4.27	3.39	5.35	7.91	6.17	5.79	5.24	4.34										
HEMFB1002502	0.83	0.8	0.28	1.27	3.14	4.39	2.38	2.95	1.77					*		+			
HEMFB1002509	0.76	0.61	0.36	0.32	0.93	0.91	0.52	1.26	0.72										
HEMFB1002518	2.29	0.9	0.49	1.25	0.69	0.67	0.59	1.16	0.95										
HEMFB1002520	10.96	4.42	7.37	13.08	19.28	16.87	8.43	9.05	9.26	*									
HEMFB1002532	2.46	1.73	4.71	2.71	2.15	2.36	2.66	2.31	4.74										
HEMFB1002537	9.87	7.21	7.79	8.36	11.1	10.55	7.47	6.16	5.86										
HEMFB1002539	7.03	2.68	3.29	3.79	4.83	3.48	4.44	3.46	4.55										
HEMFB1002531	2.36	2.37	1.2	1.94	1.74	2.82	1.39	2.3	1.35										
HEMFB1002534	4.63	2.48	3.25	4.66	8.41	8.39	2.99	3.62	3.89										
HEMFB1002536	2.96	1.03	1.7	1.05	3.49	2.9	1.99	2.14	1.93										
HEMFB1002544	3.87	12.89	3.66	4.05	4.44	5.77	1.99	5.33	2.36										
HEMFB1002545	6.5	3.17	3.97	5.87	8.72	7.62	5.47	5.22	6.78										
HEMFB1002550	3.53	1.59	2.38	1.73	2.12	4.1	3.45	2.4	2.04										
HEMFB1002556	8.37	2.84	4.27	10.84	11.6	10.64	5.58	6.3	7.41	*									
HEMFB1002571	11.52	7.77	9.15	11.56	13.65	12.93	12.05	12.33	11.31										
HEMFB1002579	9.78	5.85	5.85	7.97	13.11	12.32	6.51	5.4	6.55										
HEMFB1002582	7.48	3.22	3.33	10.72	9.33	10.11	3.01	4.39	4.41	*									
HEMFB1002584	5.81	3.4	4.16	3.75	2.97	2.76	1.46	2.06	1.93	*									
HEMFB1002587	12.23	4.61	5.2	12.45	17.92	18.78	8.13	9.27	7.5	*									
HEMFB1002590	5.23	2.47	3.26	5.42	7.78	6.92	3.77	4.99	5.06	*									

1561

1562

HEMFB1002596	11.09	4.04	5.16	6.59	10.3	10.29	7.09	7.57	6.28										
HEMFB1002600	3.89	1.64	1.46	3.06	2.9	2.86	2.13	3.4	3.88										
HEMFB1002601	4.5	1.39	1.18	5.04	4.66	4.04	3.46	3.02	3.12										
HEMFB1002603	4.45	2.06	2.73	4.75	4.46	7.27	5.07	4.62	4.32										
HEMFB1002607	3.19	2.05	1.88	4.13	5.39	8.14	1.36	2.63	2.56*										
HEMFB1002610	1.6	0.63	1.12	0.91	2.52	2.41	0.43	2.1	1.33										
HEMFB1002613	5.8	3.36	3.29	5.19	8.86	7.2	3.09	3.91	3.13										
HEMFB1002614	1.91	1.05	1.32	2.97	5.34	5.46	7.36	8.1	8.6*										
HEMFB1002615	6.52	2.3	1.68	3.51	2.94	3.31	2.18	2.84	3.84										
HEMFB1002617	2.28	1.5	2.31	5.27	5.83	5.57	3.83	2.69	3.29**										
HEMFB1002623	5.51	3.51	3.7	8.51	8.93	10.54	4.79	3.06	5.59**										
HEMFB1002624	8.23	4.59	5.1	6.42	9.16	10.04	4.11	4.54	4.4										
HEMFB1002631	1.08	1.05	0.85	1.12	1.79	1.91	1.08	2.01	0.67										
HEMFB1002635	2.64	1.42	1.61	2.73	3.71	3.6	2.53	2.71	1.37*										
HEMFB1002644	8.49	6.36	7.31	6.79	8.07	10.17	5.35	5.79	6.57										
HEMFB1002644	5.54	2.29	1.98	4.78	6.75	4.59	5.18	4.74	4.09										
HEMFB1002661	7.71	3.01	2.12	14.08	5.44	5.88	4.41	4.24	3.58										
HEMFB1002663	6.55	2.14	3.41	6.43	8.16	7.85	4.77	5.41	5.8										
HEMFB1002664	6.6	3.98	5.84	6.11	8.43	8.44	6.92	5.8	5.93										
HEMFB1002677	0.49	0.35	0.24	0.79	1.17	0.86	0.54	1.89	0.92*										
HEMFB1002683	4.48	3.9	3.87	8.9	10.99	11.79	5.35	4.88	6.92**										
HEMFB1002684	1.16	0.65	1	2.27	2.67	2.14	1.24	1.93	1.3**										
HEMFB1002686	2.67	1.11	1.21	1.17	1.78	1.98	0.85	2.28	1.79										
HEMFB1002692	1.09	0.83	0.68	1.18	2.26	3.02	1.37	1.16	1.64										
HEMFB1002693	15.96	10.15	10.49	21.46	23.57	25.74	17.35	13.97	17.93**										
HEMFB1002697	2.36	2.43	3.54	11.69	11.93	8.98	4.98	6.73	4.87**										
HEMFB1002699	13.26	6.7	7.9	16.74	17.15	20.25	11.78	11.33	10.9*										
HEMFB1002702	1.17	1.29	1.36	2.27	1.04	3.55	1.45	4.46	2.44										
HEMFB1002705	6.1	3.71	4.11	7.64	8.16	7.66	4.07	5.33	4.38*										
HEMFB1002712	1.15	0.19	1.21	2.36	1.07	1.65	1.32	2.34	0.92										
IMR321000028	14.59	7.8	9.64	7.27	7.89	8.64	3.38	5.26	3.94										
IMR321000031	3.67	1.78	1.78	4.24	3.4	4.34	3.69	3.39	3.59										
IMR321000034	26.92	15.48	13.01	18.47	24.81	26.67	19.77	14.09	22.91										
IMR321000039	17.93	8.99	10.18	11.47	11.22	20.12	13.91	11.79	14.04										
IMR321000044	0.32	0.19	0.19	0.47	1.02	1.05	0.71	2.7	0.69*										
IMR321000063	54.36	30.23	33.89	54.62	56.68	67.83	34.49	32.64	37.87										
IMR321000083	21.71	12.85	13.46	11.07	12.01	16.43	14.38	12.89	14.05										
IMR321000089	3.32	1.43	2.9	5.84	3.39	4.37	2.16	3.41	3.89										
IMR321000091	5.29	4.33	6.45	10.44	10.54	14.12	6.4	9.24	7.99**										
LIVER1000004	3.29	1.11	1.67	1.51	1.5	1.97	2.55	2.25	2.71										
LIVER1000008	3.19	0.85	0.9	1.97	1.35	1.87	1.63	1.58	2.33										
LIVER1000011	7.48	3.96	4.16	3.89	4.34	5.74	4.62	4.33	4.73										
LIVER1000022	18.53	8.45	9.73	12.74	12.74	14.58	13.3	11.15	13.79										
LIVER1000025	7.77	2.12	4.44	3.72	7.23	8.2	3.81	4.34	4.79										
LIVER1000030	4.56	1.88	1.59	2.3	3.48	3.86	1.46	2.61	2.79										
LIVER1000045	2.68	1.73	3.56	1.99	4.14	2.47	1.85	3.55	1.86										
LIVER1000046	6.12	3.71	3.54	3.3	3.9	5.04	5.21	3.87	9.2										
LIVER1000072	2.92	1.19	0.82	1.98	3.04	1.6	2.51	2.14	2.54										
LIVER1000077	4.63	3.26	3.43	3.77	4.63	3.6	5.23	4.42	4.9										
LIVER1000080	2	1.34	1.23	2.91	3.37	3.99	2.78	3.35	3.42**										
LIVER1000086	4.56	1.24	1.67	1.64	5.31	2.33	4.25	3.53	3.07										
LIVER1000092	2.68	1.43	1.4	3.38	2.77	3.26	2.88	3.19	2.12*										
LIVER1000095	4.08	1.45	1.83	2.66	3.55	3.63	2.08	3.97	1.97										
LIVER1000097	2.68	0.88	1.06	2.99	2.32	2.56	2.6	1.48	1.37										
LIVER1000098	2.82	0.74	1.66	1.13	2.25	2.13	2.82	1.76	2.99										
LIVER1000100	8.61	3.08	3.61	4.27	5.23	7.01	4.06	5.3	6.22										

43/180

【0684】

【表213】

1564

LIVER1000101	3.81	2.12	1.66	2.9	3.56	2.76	4.13	3.57	3.85				
LIVER1000106	3.32	1.56	1.67	3.52	2.18	3.06	2.2	1.66	2.75				
LIVER1000108	2.84	1.54	1.24	2.99	3.68	3.4	2.48	3.48	3.39	*	+		
LIVER1000115	2.61	1.46	1.12	3.02	3.28	3.44	1.96	2.92	2.86	*	+		
LIVER1000120	5.02	2.94	2.41	3.82	3	3.25	3.35	2.12	2.66				
LIVER1000138	4.91	0.99	2.36	1.52	2.93	3.2	2.89	4.6	2.68				
LIVER1000146	11.83	5.09	5.8	8.13	11.73	11.21	7.01	6.1	7.3				
LIVER1000148	11.43	4.5	7.19	7.38	7.37	7.43	6.46	5.27	6.13				
LIVER1000157	33.53	16.69	18.55	25.58	33.97	31.92	16.84	15.36	18.47				
LIVER1000161	7.22	4.95	3.61	5.26	5.68	6.24	4.45	3.94	6.08				
LIVER1000167	4.56	2.13	2.81	3.19	3.07	3.13	1.51	2.38	2.42				
LIVER1000174	3.84	1.31	1.5	1.69	2.19	2.47	1.08	2.69	2.65				
LIVER1000185	6.12	3.35	4.22	3.51	3.56	2.98	2.75	3.21	2.98				
LIVER1000187	3.26	1.56	0.93	1	1.39	1.74	0.82	3.36	0.61				
LIVER1000190	1.93	1.11	1.59	1.96	1.59	2.03	2.89	1.41	1.66				
LIVER1000192	10.65	6.24	5.2	5.75	5.77	6.49	6.06	5.8	6.02				
MAMMA1000009	5.3	2.68	2.46	6.62	5.77	8.83	4.6	3.23	5.23	*	+		
MAMMA1000015	5.84	1.77	1.87	1.44	2.88	3.59	3.38	2.5	2.77				
MAMMA1000019	5.66	2.6	2.84	4.89	9.82	8.95	3.81	3.64	4.85				
MAMMA1000020	3.8	3.44	4.09	3.56	8.72	8.06	4.37	4.09	4.16				
MAMMA1000024	2.87	0.82	0.95	1.1	1.88	2.53	1.55	2.13	2.01				
MAMMA1000025	4.87	2.19	2.6	4.8	5.71	6.47	3.27	3.34	4.13				
MAMMA1000043	10.51	5.09	5.02	14.31	20.26	13.23	7.72	9.62	9.43	*	+		
MAMMA1000046	1.69	0.97	1.62	2.91	3.36	3.57	3.47	1.81	3.55	**	+		
MAMMA1000046	6.47	2.08	3.57	6.03	7.6	8.45	5.17	3.75	4.66				
MAMMA1000055	6	3.15	3.53	2.8	3.48	4.97	5.81	4.07	2.35				
MAMMA1000057	12.48	5.52	7.03	12.15	20.3	15.59	7.03	7.1	8.26				
MAMMA1000060	14.45	7.18	9.91	16.29	13.21	18.23	10.59	9.1	11.91				
MAMMA1000069	7.73	3.61	4.66	6.69	8.82	10.74	4.08	5.81	4.8				
MAMMA1000084	9.73	3.57	5.05	11.91	14.34	16.88	5.45	7.65	6.73	*	+		
MAMMA1000085	3.47	1.96	1.87	2.74	2.35	3.06	1.99	2.32	2.6				
MAMMA1000092	5.41	2.13	2.26	4.85	6.6	6.02	2.97	4.24	4.71				
MAMMA1000096	3.78	3.03	1.78	3.72	4.8	6.47	4.17	3.9	6.06				
MAMMA1000097	5.13	2.95	3.91	5.32	4.24	6.86	3.6	3.62	3.69				
MAMMA1000102	4.12	2.21	2.7	5.22	5.81	5.02	2.36	4.65	3.65				
MAMMA1000103	3.31	1.56	2.28	4.58	6.05	6.54	2.94	2.99	3.37	*	+		
MAMMA1000106	2.7	1.79	2.13	3.04	5.09	5.41	1.36	3.69	2.47	*	+		
MAMMA1000117	2.72	1.52	1.22	1.31	2.51	2.71	0.5	1.62	1.27				
MAMMA1000118	8.14	2.71	2.77	3.78	7.64	6.37	5.72	5.22	4.29				
MAMMA1000129	4.52	1.62	2.67	3.35	3.9	5.18	1.94	2.89	2.82				
MAMMA1000133	4.27	1.92	2.22	2.89	3.17	3.71	2.86	2.72	3.28				
MAMMA1000134	3.24	1.82	3.24	6.48	6.88	8.35	3.29	3.76	4.59	**	+		
MAMMA1000139	3.29	2.4	1.31	3.92	4.25	4.14	3.22	2.8	2.68	*	+		
MAMMA1000141	3.46	1.27	2.24	4.07	4.79	6.79	1.97	2.52	1.91	*	+		
MAMMA1000143	2.16	0.91	1.71	2.99	2.74	3.39	1.31	2.55	1.46	*	+		
MAMMA1000159	10.88	7.04	8	8.79	14.06	12.93	3.84	10.55	5.74				
MAMMA1000158	10.85	5.54	5.47	9.19	13.85	13.81	7.6	7.75	9.58				
MAMMA1000163	5.58	3.38	2.67	5.07	6.46	5	2.15	2.84	6.5				
MAMMA1000171	7.29	4.5	8.08	8.95	12.01	14.93	6.64	7.02	7.82	*	+		
MAMMA1000173	6.86	4.32	5.72	5.71	7.66	7.6	5.97	5.63	5.95				
MAMMA1000175	4.12	1.18	0.23	1.4	1.36	1.53	1.51	3.3	1.81				
MAMMA1000183	7	6.5	5.17	8.02	15.13	12.18	4.61	6.16	5.98				
MAMMA1000191	6.82	3.67	4.83	4.54	6	5.86	3.61	5.7	4.78				
MAMMA1000192	13.21	7.3	7.84	8.79	11.31	9.83	5.1	9.07	11.26				
MAMMA1000193	6.03	2.64	1.36	3.73	7.78	4.43	3.35	3.38	4.25				
MAMMA1000196	11.19	3.7	4.24	11.67	15.53	12.34	7.74	5.71	7.75				

44/180

【表 2 1 4】

1565

1566

MAMMA1000284	7.62	4.33	5.82	9.51	6.8	8.34	5.56	4.69	5.2										
MAMMA1000287	6.14	2.58	4.15	4.25	4.43	7.49	4.2	4.63	3.58										
MAMMA1000214	3.73	2.36	3.5	6.05	6.43	8.36	3.87	5.19	4.02*	+									
MAMMA1000220	3.64	2.49	2.27	4.02	3.64	4.91	4.36	4.51	3.83									*	+
MAMMA1000221	4.11	1.84	1.12	2.33	12.39	3.34	2.86	3.69	1.65										
MAMMA1000226	3.4	1.09	2.76	2.96	2.31	2.84	1.92	4.54	2.53										
MAMMA1000227	5.88	3.58	3.47	4.08	7.55	8.07	3.93	3.56	5.9										
MAMMA1000230	6.36	3.63	3.36	3.79	7.14	7.18	4.32	4.39	3.89										
MAMMA1000241	5.23	2.78	2.92	6.17	10.99	9.16	5.63	6.94	5.92*	+	*	+							
MAMMA1000245	71.79	48.41	41.99	49.62	55.47	70.51	36.86	32.29	42.56										
MAMMA1000248	10.75	5.11	8.19	10.32	13.93	13.73	8.64	7.83	9.87										
MAMMA1000251	4.47	3.42	3.86	6.07	8.71	10	3.62	6.05	5.41*	+									
MAMMA1000254	2.89	1.15	1.35	4.06	3.79	5.07	1.95	3.72	2.71*	+									
MAMMA1000257	7.12	4.26	6.71	11.96	14.47	16.44	5.81	9.74	10.27**	+									
MAMMA1000262	12.13	6.11	6.35	9.28	17.3	14.89	11.45	12.94	13.68										
MAMMA1000264	1.54	1.94	1.06	2.96	5.16	6.26	1.9	2.25	1.92*	+									
MAMMA1000266	1.41	0.76	1.44	2.49	3.39	2.45	2.4	2.54	1.43*	+									
MAMMA1000270	8.33	3.85	6.34	9.35	14.72	13.36	5.23	6.87	8.24*	+									
MAMMA1000271	3.79	2.55	1.83	6.46	5.81	4.43	3.8	4.01	4.5*	+									
MAMMA1000277	2.17	1.07	1.86	2.66	2.04	3.91	1.48	2.33	1.37										
MAMMA1000278	2.46	1.53	1.53	2.26	1.74	1.78	1.61	3.39	1.57										
MAMMA1000279	4.53	3.12	3.68	7.71	9.92	13.85	2.86	4.21	4.62*	+									
MAMMA1000283	2.8	0.74	1.34	2.2	3.06	3.24	2.27	2.64	2.53										
MAMMA1000284	7.09	3.1	3.89	5.31	5.61	7.3	4.33	4.12	6.21										
MAMMA1000287	3.34	1.37	2.39	5.26	5.17	6.99	4.97	3.06	4.33*	+									
MAMMA1000294	18.13	8.47	8.55	15.53	11.48	16.82	12.33	10.64	11.59										
MAMMA1000298	1.54	0.71	0.82	0.74	1.91	1.79	1.37	1.29	1.02										
MAMMA1000302	5.12	2.71	2.69	5.15	5.37	6.89	4.36	4.77	2.99										
MAMMA1000303	4	2.05	1.59	2.54	3.44	3.95	1.95	2.67	2.43										
MAMMA1000305	1.38	0.71	0.71	1.7	2.67	3.22	1.16	1.69	1.13*	+									
MAMMA1000307	12.76	5.57	7.52	10.78	17.15	13.46	11.84	12.09	11.6										
MAMMA1000309	0.76	0.89	1.4	1.06	1.34	1.72	1.77	0.93	1.2										
MAMMA1000312	1.8	1.04	0.87	1.28	0.56	1.1	1.25	1.47	0.9										
MAMMA1000313	2.67	3.77	1.89	3.1	6.23	5.66	3.12	2.28	2.98										
MAMMA1000331	4.12	2.28	1.93	3.93	3.97	5.29	3.56	3.45	3.82										
MAMMA1000335	6.16	2.7	3.37	3.54	3.79	3.88	3.68	2.45	3.73										
MAMMA1000339	3.25	1.33	2.61	3.01	4.9	3.33	2.91	2.77	1.92										
MAMMA1000340	2.6	1.63	1.41	3.96	4.43	4.29	1.81	3.28	2.22**	+									
MAMMA1000348	3.33	1.48	2.34	6.45	6.9	6.21	5.1	3.51	6.66**	+									
MAMMA1000356	8.13	2.7	3.74	9.76	8.55	10.65	5.97	5.34	5.67										
MAMMA1000358	4.37	2.17	1.44	5.1	4.35	4.38	3.5	3.09	3.71										
MAMMA1000360	7.72	3.05	2.69	11.41	9.78	10.42	6.57	4.42	6.39*	+									
MAMMA1000361	7.91	2.97	4.89	10.45	10.37	13.01	6.44	5.43	7.13*	+									
MAMMA1000363	5.44	2.67	2.71	3.44	2.89	4.74	2.99	2.83	3.16										
MAMMA1000370	8.4	6.64	6.2	6.19	7.25	6.56	6.68	7.49	4.91										
MAMMA1000371	6.81	4.41	6.08	4.39	3.58	5.6	4.96	6.77	5.24										
MAMMA1000372	11.86	4.03	5.88	15.22	16.38	16.77	7.36	6	7.47*	+									
MAMMA1000385	4.62	2.3	2.77	5.18	7.04	8.05	4.85	4.48	5*	+									
MAMMA1000388	6.44	2.83	3.67	5.65	4.46	4.85	4.91	3.34	5.06										
MAMMA1000395	5.17	2.17	2.95	3.65	4.16	4.78	3.21	2.41	3.84										
MAMMA1000403	7.68	3.41	2.88	9.51	10.11	10.62	5.46	6.68	5.96*	+									
MAMMA1000403	6.72	2.73	3.78	6.04	7.7	8.56	4.71	5.83	4.03										
MAMMA1000410	4.02	2.21	1.56	4.09	5.7	5.12	2.32	3.4	1.98										
MAMMA1000413	1.97	0.9	1.1	2.1	2.16	1.61	0.81	7.47	1.21										
MAMMA1000414	3.35	1.71	2.96	4.73	3.34	2.27	4.52	4.04	1.89										
MAMMA1000416	14.38	8.87	8.86	11.04	19.59	15.46	12.54	10.99	12.32										

45/180

1568

MAMMA1000421	7.88	5.58	3.16	7.31	11.57	11.97	5.34	5.28	5.81								
MAMMA1000422	4.93	2.59	1.84	2.34	3.07	4.44	2.35	2.99	4.14								
MAMMA1000423	3.67	2.68	1.35	2.17	3.71	4.12	2.5	2.73	2.24								
MAMMA1000424	0.47	0.75	0.45	1.27	1.57	1.76	1.14	1.64	1.04	**	+	*					
MAMMA1000429	32.94	14.89	22.85	23.37	29.25	34.16	20.87	25.24	25.81								
MAMMA1000431	7.98	3.3	4.81	7.45	10.34	10.08	4.8	6.35	5.12								
MAMMA1000432	4.6	2.09	3.06	2.28	3.64	3.72	3.49	4.61	3.15								
MAMMA1000437	6.14	5.61	6.7	6.37	13.88	6.85	9.07	7.74	6.38								
MAMMA1000444	10.06	5.02	5.92	12.4	21.04	12.87	7.66	9.16	10.21								
MAMMA1000446	5.86	2.32	2.37	3.48	5.41	5.04	2.92	3.11	3.2								
MAMMA1000449	5.06	1.88	4.07	4.87	7.02	6.19	3.35	3.99	3.47								
MAMMA1000457	3.42	1.31	1.57	3.54	3.24	3.66	3.14	3.48	3.29								
MAMMA1000459	3.87	1.25	2.08	2.19	3.1	2.93	2.24	2.82	1.85								
MAMMA1000469	1.49	0.06	0.79	0.79	1.06	1.13	0.34	1.08	0.62								
MAMMA1000472	11.38	4.74	6.91	9.53	12.61	11.92	6.13	7.53	8.61								
MAMMA1000473	5.96	3.57	3.53	12.63	7.19	13.81	5.26	5.18	5.28	*	+	*					
MAMMA1000477	5.82	2.74	2.51	5.72	8.15	7.58	3.74	4.02	3.75								
MAMMA1000478	9	4.17	4.73	12.94	18.52	17.59	8.49	7.88	8.95	*	+	*					
MAMMA1000483	14.86	5.67	8.42	11.14	12.63	12.05	7.76	6.25	5.28								
MAMMA1000499	3.41	1.2	1.17	3.21	2.92	3.1	1.71	2.32	2.64								
MAMMA1000496	2.46	1.87	1.02	2.44	3.29	2.49	1.44	3.16	1.85								
MAMMA1000509	1.56	0.84	0.9	2.28	2.73	1.98	1.08	1.9	1.36	*	+	*					
MAMMA1000501	11.66	5.38	5.27	11.85	14.49	13.05	6.88	6.5	10.43								
MAMMA1000503	1.33	0.54	0.92	1.59	1.74	1.27	1.8	2.54	1.09								
MAMMA1000506	12.82	9.48	10.39	12.58	12.2	12.4	9.73	8.88	12.24								
MAMMA1000510	7.01	5.28	6.34	4.55	7.48	6.53	5.31	5.02	4.6								
MAMMA1000518	7.48	2.78	3.25	5.65	6.45	7.72	3.13	3.76	3.48								
MAMMA1000516	5.84	1.59	2.82	5.98	7.85	7.2	2.82	3.57	3.21								
MAMMA1000522	2.27	1.18	1.41	3.64	3.92	3.54	1.42	3.62	1.52	**	+	*					
MAMMA1000524	7.63	2.43	4.92	8.34	11.81	13.33	5.04	5.34	4.54	*	+	*					
MAMMA1000528	1.65	0.58	1.07	2.05	2.46	2.53	1.6	1.39	1.82	*	+	*					
MAMMA1000534	2.5	1.5	1.3	2.79	2.83	2.9	2.6	2.21	1.6	*	+	*					
MAMMA1000541	10.98	5.23	5.03	6.32	9.31	8.45	6.48	6.33	7.6								
MAMMA1000550	4.4	3.04	2.74	4.35	5.4	3.92	4.73	3.37	2.94								
MAMMA1000556	1.48	1.03	1.14	1.83	2.63	2.37	0.93	2.78	1.93	*	+	*					
MAMMA1000559	4.37	1.96	1.73	4.8	7.23	5.02	4.99	3.84	3.11								
MAMMA1000566	4.72	1.49	2.86	6.83	6.65	5.82	4.27	3.68	2.63	*	+	*					
MAMMA1000567	3.83	3.37	3.67	5.22	7.17	6.61	3.18	4.82	3.63	*	+	*					
MAMMA1000576	15.99	9.01	6.07	17.14	30.24	29.01	12.9	10.14	12.06	*	+	*					
MAMMA1000582	5.54	2.74	3.08	4.19	5.56	6.62	5.53	3.7	3.87								
MAMMA1000583	4.38	2.28	1.5	5.07	4.73	6.13	4.1	3.32	3.54								
MAMMA1000585	3.99	1.32	2.85	5.97	7.85	8.52	3.94	4.82	4.06	*	+	*					
MAMMA1000587	3.21	2.47	2	4.38	5.07	2.06	5.51	4.86	2.27								
MAMMA1000591	3.28	1.11	2.12	2.42	2.51	3.46	1.69	4.06	3.09								
MAMMA1000594	6.52	3.99	5.77	13.18	11.99	15.24	6.18	7.35	5.58	**	+	*					
MAMMA1000597	17.18	8.64	13.27	24.68	32.8	31.71	15.42	12.85	17.87	*	+	*					
MAMMA1000605	15	7.83	7.51	16.1	27.17	27.84	15.98	12.24	13.96	*	+	*					
MAMMA1000612	7.9	2.22	3.52	4.29	4.53	4.73	3.74	1.84	4.78								
MAMMA1000614	21.9	15.16	16.51	11.47	18.81	18.51	16.91	14.24	15.41								
MAMMA1000616	0.69	0.1	0.08	2.78	1.16	2.29	1.88	2.45	1.31	*	+	*					
MAMMA1000621	3.29	2.06	2.49	3.72	4.74	3.92	2.54	4.56	2.58								
MAMMA1000623	3.66	0.62	3.18	3.6	1.78	3.6	1.68	2.93	1.92								
MAMMA1000625	21.85	13.69	19.79	20.91	20.47	21.13	14.29	16.77	23.93								
MAMMA1000635	0.42	0.29	0.29	1.14	0.49	1.07	0.77	1.98	0.45								
MAMMA1000640	3.78	2.57	1.76	3.32	6.22	6.75	3.82	3.69	4.44	*	+	*					
MAMMA1000646	10.28	5.04	4.34	5.25	6.7	11.98	4.93	10.71	4.80								

46/180

【表 2 1 6】

1569

1570

МАММА1008652	8.47	3.81	3.01	8.32	13.85	13.09	5.34	6.27	6.14	*	+				
МАММА1008657	5.07	3.94	3.85	6.77	10.59	9.73	5.63	6.14	5.11	*	+				
МАММА1008664	2.69	1.1	1.96	4.49	4.5	4.2	5.68	4.29	2.35	**	+				
МАММА1008667	4.79	1.98	2.15	4.21	4.93	5.76	3.08	4.06	3.71	*	+				
МАММА1008669	2.4	1.13	1.67	3.73	2.97	3.09	0.95	4.13	2.08	*	+				
МАММА1008669	1.17	0.4	0.79	2.08	2.59	2.37	1.24	0.92	0.96	**	+				
МАММА1008670	7.56	4.44	3.7	4.32	4.44	6.75	2.59	5.1	5.48	*	+				
МАММА1008672	7.79	2.99	3.4	4.22	3.53	6.53	3.72	4.19	6.43	*	+				
МАММА1008681	4.68	1.14	3.03	2.41	2.85	4.06	2.7	2.22	3.58	*	+				
МАММА1008684	35.85	22.61	24.91	21.42	31.5	29.68	12.4	13.65	15.36	*	+				
МАММА1008694	6.4	3.52	4.51	7.83	11.25	15.25	8.55	6.27	7.54	*	+				
МАММА1008702	8.51	4.05	5.46	6.26	5.22	7.23	5.02	5.02	4.55	*	+				
МАММА1008706	3.68	1.19	1.86	2.9	2.36	3.42	1.81	1.88	2.14	*	+				
МАММА1008707	3.62	1.77	1.28	1.62	3.45	1.98	2.41	2.52	2.5	*	+				
МАММА1008713	5.4	2.54	3.24	5.36	5.73	6.33	4.52	4.76	4.87	*	+				
МАММА1008714	7.46	4.12	5.15	8.57	7.81	8.68	8.73	7.85	8.07	*	+				
МАММА1008718	3.29	2.59	1.62	6.31	6.72	5.21	3.55	3.17	4.84	**	+				
МАММА1008720	11.1	3.49	5.25	10.45	13.49	12.85	6.43	5.97	7.74	*	+				
МАММА1008723	2.28	1.69	2.12	4.14	3.59	4.23	2.79	2.97	1.93	**	+				
МАММА1008731	1.86	0.62	0.69	2.69	3.19	3.37	2.54	2.31	2.78	*	+	*	+		
МАММА1008732	4.46	2.1	1.55	3.27	6.08	6	3.73	4.07	3.22	*	+				
МАММА1008733	2	0.47	0.64	1.76	2.5	2.33	0.99	1.71	0.61	*	+				
МАММА1008734	19.84	13.3	8.71	14.98	15.8	18.61	13.99	14.24	10.98	*	+				
МАММА1008736	12.43	4.93	6.22	7.65	6.62	9.44	6.16	4.05	8.82	*	+				
МАММА1008738	9.86	3.76	4.66	5.29	7.95	8.71	4.04	5.76	4.24	*	+				
МАММА1008744	6.53	4.63	4.71	11	10.23	11.31	6.26	6.39	7.79	**	+				
МАММА1008746	1.48	2.11	1.07	4.85	6.59	5.04	2.55	4.44	6.26	**	+				
МАММА1008748	9.39	7.13	8.61	8.38	10.56	16.11	5.63	9.36	9.45	*	+				
МАММА1008751	19.32	15.21	15.9	12.13	17.33	24.65	8.32	12.47	10.06	*	-				
МАММА1008752	4.99	3.06	2.62	6.31	5.93	7.52	3.37	3.3	3.21	*	+				
МАММА1008757	16.42	7.46	8.63	15.03	20.13	20.42	10.82	9.38	12.45	*	+				
МАММА1008760	13.83	4.85	6.07	16.93	20.12	21.36	9.26	10.09	9.12	*	+				
МАММА1008761	7	5.05	5.28	10.4	11.63	13.03	5.96	6.75	6.32	**	+				
МАММА1008775	4.08	1.66	2.88	3.15	4.48	7.4	3.92	4.45	3.2	*	+				
МАММА1008776	6.7	4.59	3.36	9.35	9.08	9.79	6.68	5.65	5.84	*	+				
МАММА1008778	5.98	3.45	2.59	7.46	6.58	10.39	4.17	4.73	3.98	*	+				
МАММА1008781	5.48	3.83	3.81	4.84	4.93	5.96	2.78	5.06	3.06	*	+				
МАММА1008783	15.43	7.59	9.38	7	8.75	12.93	6.89	10.66	10.04	*	+				
МАММА1008784	6.69	3.02	3.41	4.23	8.26	6.49	8.78	3.6	3.47	*	+				
МАММА1008788	18.64	7.23	10.16	10.93	9.2	11.24	9.78	6.25	8.61	*	+				
МАММА1008798	2.84	1.31	1.29	2.57	6.45	2.47	2.42	2.49	2.05	*	+				
МАММА1008803	10.19	4.79	5.55	11.64	14.85	12.54	8.45	6.23	7.37	*	+				
МАММА1008810	10.4	8.43	5.83	11.45	14.19	14.79	8.3	8.84	9.48	*	+				
МАММА1008813	3.06	1.41	1.3	0.97	1.08	1.47	1.17	2.87	1.61	*	+				
МАММА1008814	11.43	4.36	6.48	10.9	13.12	14.78	6.64	8.56	8.44	*	+				
МАММА1008824	4.94	1.4	2.5	6.51	8.16	10.38	6.57	7.55	6.92	*	+	*	+		
МАММА1008827	5.81	3.08	3.37	6.5	5.83	6.58	3.91	3.77	4.74	*	+				
МАММА1008831	3.49	2.19	2.43	2.04	2.83	2.54	2.49	2.54	3.51	*	+				
МАММА1008838	7.72	3.74	6.73	10.55	7.02	15.37	8.46	6.62	9.4	*	+				
МАММА1008839	9.86	5.11	5.3	13.32	14.94	15.98	11.39	9.61	11.68	**	+				
МАММА1008841	2.16	2.22	2.46	2.34	3.62	2.61	2.07	3.51	3.1	*	+				
МАММА1008843	9.7	5.15	5.18	5.26	8.56	8.54	4.59	6.92	6.8	*	+				
МАММА1008843	1.45	0.52	0.63	1.44	1.42	1.66	1.24	1.97	1.12	*	+				
МАММА1008845	2.99	0.85	1.74	2.17	3.73	3.02	1.45	3.21	2.14	*	+				
МАММА1008851	12.84	5.8	5.26	10.17	14.4	13.52	7.61	8.15	8.58	*	+				
МАММА1008854	5.64	2.1	2.3	3.44	3.33	5.81	6.81	5.97	6.68	*	+				

47/180

1572

MAMMA100085	1.7	1.63	1.03	1.59	2.99	3.96	1.06	2.13	1.04						
MAMMA100086	6.3	3.91	3.68	6.66	6.53	6.39	5.69	5.47	5.67						
MAMMA100089	30.54	14.5	21.77	15.43	16.32	21.44	10.77	8.93	11.82						
MAMMA100090	3.63	1.84	2.53	2.21	2.9	4.05	1.42	1.82	1.19						
MAMMA100093	6.2	3.01	3.04	4.59	9.69	8.1	4.1	6.66	5.5						
MAMMA100096	0.8	0.11	0.15	0.67	1.37	0.92	0.2	1.71	0.5						
MAMMA100097	4.15	2.15	1.95	2.19	5.49	3.51	1.75	2.5	2.37						
MAMMA1000975	9.92	4.24	6.11	6.91	11.92	12.78	4.67	4.48	7						
MAMMA1000976	4.63	2.26	3.14	3.33	5.28	6.68	4.51	3.48	5.36						
MAMMA1000977	9.58	4.24	6.31	9.18	13.08	15.47	7.32	6.45	8.51						
MAMMA1000978	8.16	4.46	5.1	7.91	13.1	10.3	5.72	5.68	6.98						
MAMMA100098	4.25	2.2	2.38	4.84	4.93	5.5	2.27	3.49	2.89	*	+				
MAMMA1000981	4.86	3.39	4.01	5.58	9.07	9.97	3.7	4.59	4.69	*	+				
MAMMA1000983	4.1	2.09	3.9	3.29	3.78	3.16	2.41	3.12	3.57						
MAMMA1000997	0.87	0.78	1.52	1.35	2.84	1.6	1.61	1.81	0.9						
MAMMA1000998	14.3	5.37	5.9	6.61	8.53	8.24	7.24	7.58	9.2						
MAMMA1000995	6.32	4.18	3.03	7.58	8.06	10.95	4.06	4.04	6.22	*	+				
MAMMA1000996	4.24	2.45	3	4.3	3.89	5.72	2.87	4.2	3.18						
MAMMA1000999	1.37	0.39	0.86	1.42	2.93	1.74	2.49	2.77	1.87			*	+		
MAMMA10009911	0.61	1.25	0.84	1.86	2.28	6.33	8.08	5.76	7.77	*	+	**	+		
MAMMA10009914	5.03	2.41	2.68	4.67	4.17	3.32	1.99	2.14	2.33						
MAMMA10009920	3.12	1.17	2.51	3.63	3.17	3.45	2.05	3.06	3.19						
MAMMA10009921	3.37	3.29	3.26	3.61	9.57	6.95	3.48	3.25	3.54						
MAMMA10009931	8.02	4.92	5.62	10.36	14.6	15.07	6.35	6.66	5.94	*	+				
MAMMA10009940	6.43	3.57	4.1	8.17	7.42	11.2	5.43	7.24	5.59	*	+				
MAMMA10009941	8.08	4.42	5.26	11.96	15.08	14.97	7.8	6.29	5.57	**	+				
MAMMA10009943	16.28	7.28	9.32	16.51	16.66	17.99	9.16	10.49	11.15						
MAMMA10009949	8.02	5.62	7.75	12.59	16.34	17.28	9.76	9.93	7.72	**	+				
MAMMA10009952	8.49	4.92	6.82	13.66	13.4	12.11	7.68	8.43	10.02	**	+				
MAMMA10009954	1.29	1.15	1.49	1.35	3.18	2.29	2.16	3.08	2.19			*	+		
MAMMA10009957	6.37	3.36	2.47	7.39	11.27	10.47	4.72	6	5.03	*	+				
MAMMA10009962	14.04	6.88	6.94	17.04	23.21	26.2	11.63	8.86	12.79	*	+				
MAMMA10009966	7.34	3.73	4.5	10.84	15.74	12.34	4.66	6.62	6			*	+		
MAMMA10009968	7.71	3.48	2.83	8.85	11.98	9.01	6.3	7.27	5.97	*	+				
MAMMA10009972	1.58	1.55	1.15	3.48	2.9	3.02	2.22	4.51	2.3	*	+				
MAMMA10009973	3.5	1.69	1.59	3.69	3.21	4.33	2.55	2.9	1.2						
MAMMA10009975	2.22	2.8	2.6	2.48	6.62	3.03	2.24	4.33	2.06						
MAMMA10009976	7.5	4.17	5.75	10.05	14.48	15.04	6.28	7.31	7.44	*	+				
MAMMA10009979	6.1	3.13	2.84	6.83	11.15	7.34	4.03	3.36	5.99						
MAMMA10009986	8.92	4.73	5.33	9.12	17.71	11.66	6.36	10.27	8.03						
MAMMA10009987	4.61	3.28	2.96	7.53	9.04	9.57	3.67	3.24	4.14	**	+				
MAMMA10009988	6.9	4.02	3.13	9.98	9.41	10.85	6.42	4.87	6.36	*	+				
MAMMA10009994	3.37	2.44	3.14	3.13	4.33	4.9	3.61	4.21	3.97			*	+		
MAMMA10009999	3.52	2.26	2.81	4.12	6.42	7.42	3.48	4.56	3.6	*	+				
MAMMA10010003	1.84	1.4	1.47	5.67	6.98	6.99	2.14	3.71	2.23	**	+				
MAMMA10010007	0.12	0.01	0.3	0.22	0.03	0.58	0.25	0.21	0.73						
MAMMA10010088	6.4	6.37	4.3	6.99	5.97	6.01	5.02	5.89	7.81						
MAMMA1001013	6.8	3.38	4.83	15.25	11.23	8.98	5.96	5.59	9.13	*	+				
MAMMA1001014	7.78	3.67	2.44	4.42	6.29	6.7	2.43	2.82	2.35						
MAMMA1001021	7.09	2.52	2.8	7.68	6.46	6.9	4.64	3.79	3.74						
MAMMA1001024	8.72	3.44	3.61	8.02	10.11	9.18	4.3	6.16	5.88						
MAMMA1001025	1.98	1.65	0.42	0.75	1.1	1.07	0.62	0.65	0.73						
MAMMA1001028	3.61	3.77	2.41	1.41	2.09	2.3	1.65	2.01	1.32	*	-				
MAMMA1001030	3.45	1.67	2.14	3.47	2.37	4.44	0.07	2.57	2.47						
MAMMA1001035	13.14	7.47	7.89	19	23.71	18.79	11.21	8.37	14.57	**	+				
MAMMA1001036	11.51	6.94	5.48	11.14	14.27	13.18	7.47	5.06	7.52						

1573

1574

MAMMA1001037	9.85	4.28	3.71	10.53	13.73	9.2	7.98	5.87	7.42	*	*	*
MAMMA1001038	3.03	1.45	2.07	4.49	7.26	6.93	4.49	3.88	6.41	*	*	*
MAMMA1001041	6.12	4.31	3.78	4.25	5.32	5.37	5.37	4.53	3.75			
MAMMA1001043	9.46	4.63	3.66	4.68	7.75	7.15	4.92	3.72	4.88			
MAMMA1001050	6.35	5.89	3.9	5.29	10.13	10.16	5.02	6.56	5.49			
MAMMA1001054	5.51	4.13	3	8.5	8.45	8.15	5.21	3.63	4.46	**	+	
MAMMA1001059	15.39	8.08	6.23	9.1	11.74	11.86	8.44	7.77	9.49			
MAMMA1001066	16.43	8.7	9.35	16.38	15.95	15.31	10.1	8.21	12.62			
MAMMA1001067	3.67	2.44	1.56	5.04	5.4	5.91	3.35	3.05	4.31	*	+	
MAMMA1001073	11.88	5.32	6.63	6.72	4.61	6.46	5.54	4.86	5.86			
MAMMA1001073	5.21	2.94	1.75	2.04	3.72	2.45	1.94	3.03	2.39			
MAMMA1001074	3.99	4.38	2.77	4.13	9.96	7.19	3.27	3.81	5.24			
MAMMA1001075	5.54	2.96	3.2	3.06	7.9	7.5	2.62	3.18	3.18			
MAMMA1001078	7.94	4.65	4.05	9.11	13.65	11.41	7.34	5.68	7.64	*		
MAMMA1001080	22.36	9.18	10.44	11.87	12.56	12.61	9.96	10.53	13.83			
MAMMA1001082	4.52	3.3	1.66	3.03	5.82	3.36	3.3	2.6	2.45			
MAMMA1001091	0.73	0.99	0.34	1.07	1.55	1.04	1.3	1.27	1.5	*	+	*
MAMMA1001092	3.38	1.71	1.14	4.68	5.06	3.84	2.72	2.57	3.2	*	+	*
MAMMA1001094	23.07	10.75	8.74	19.47	15.51	11.95	11.1	12.09	9.06			
MAMMA1001105	8.97	7.82	3.9	7.84	13.25	10.97	5.27	6.89	7.2			
MAMMA1001110	1.34	0.28	1.07	0.83	1.4	1.91	0.64	1.83	0.87			
MAMMA1001126	11.76	5.19	6.22	18.27	20.42	20.62	10.8	7.93	10.63	**	+	
MAMMA1001133	13.96	7.98	6.29	17.52	21.82	18.6	12.41	9.09	11.57	*	+	
MAMMA1001139	16	10.86	8	75.48	52.51	50.41	4.72	2.94	4.09	**	+	*
MAMMA1001141	3.54	2.73	2.73	3.35	3.24	4.02	3.72	4.28	4.25			
MAMMA1001143	9.1	5.11	2.81	6.09	8.1	8.79	3.24	3.97	7.09			
MAMMA1001146	8.33	4.95	3.62	3.46	6.81	6.75	3.46	5.11	7.05			
MAMMA1001150	8.4	3.25	2.79	2.57	3.1	4.61	3.41	4.01	4.33			
MAMMA1001154	10.09	4.99	5.59	11.85	11.71	18.3	6.93	7.19	6.3			
MAMMA1001159	9.34	6.32	4.92	5.06	4.86	4.07	3.31	2.7	4.01			
MAMMA1001161	14.59	7.23	8.28	17.47	24.12	19.35	11.34	7.11	8.84	*	+	
MAMMA1001163	8.3	3.74	4.22	6.24	6.6	5.21	4.88	5.43	5.84			
MAMMA1001181	5.83	2.22	1.87	4.38	4.79	3.53	3.65	3.3	3.3			
MAMMA1001186	7.43	2.73	2.8	9.55	11.46	10.04	5.94	5.12	6.23	*	+	
MAMMA1001189	5.2	2.45	3.28	2.31	6.23	8.54	2.7	3.48	4.97			
MAMMA1001191	7.35	3.89	3.31	3.72	5.24	6.78	3.27	4.86	5.76			
MAMMA1001198	420.1	187.9	245.8	305.4	416.1	499.3	169.9	199.8	188.3			
MAMMA1001203	22.54	12.72	10.05	25.35	28.4	25.81	14.74	13.68	16.11	*	+	
MAMMA1001203	10.49	4.64	4.15	9.25	14.44	10.45	6.11	7.56	8.28			
MAMMA1001206	6.15	2.67	2.33	5.52	7.44	5.57	3.53	2.86	3.88	*	+	
MAMMA1001208	6.57	2.81	3.7	5.42	5.59	5.39	4.2	3.8	4.35			
MAMMA1001215	10.79	5.58	5.27	10.75	14.22	15.01	5.67	7.42	7.48			
MAMMA1001220	9.93	5.68	4.3	14.63	18.62	17.06	7.53	7.5	9.1	**	+	
MAMMA1001223	1.59	0.92	0.2	0.96	1.98	1.96	0.04	1.13	0.9			
MAMMA1001223	4.89	1.72	1.83	2.87	4.51	4.18	2.3	4.01	2.37			
MAMMA1001233	8.78	2.9	3.18	7.54	10.45	9.18	4.93	4.96	6.51			
MAMMA1001234	7.4	4.59	2.41	6.32	6.84	8.88	3.78	3.73	6			
MAMMA1001237	2.49	1.76	1.72	1.22	2.15	2.66	1.99	1.8	2.56			
MAMMA1001243	2.36	1.9	1.62	4.41	7.15	5.32	4.22	3.46	4.95	*	**	+
MAMMA1001244	2.4	1.42	0.68	1.53	3.03	2.44	2.05	2.71	3.6			
MAMMA1001249	5.06	0.96	1.74	3.77	9.25	4.93	2.44	3.57	4.08			
MAMMA1001256	2.41	7.77	2.44	2.13	6.99	4.49	2.44	2.38	3.83			
MAMMA1001259	5.56	2.92	3.02	4.36	6.71	5.33	2.23	3.99	5.03			
MAMMA1001260	13.79	6.11	6.31	13.52	13.26	12.23	7.61	6.68	10.66			
MAMMA1001263	9.64	6.71	5.97	8.72	6.36	5.96	5.41	6.02	8.33			
MAMMA1001268	4.72	2.75	3.01	9.39	6.71	7.53	3.74	4.95	5.14	*	+	*

49/180

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【表 2 1 9】

1576

MAMMA1001371	18.48	7.38	8.91	10.48	14.14	10.31	9.58	8.4	12.04				
MAMMA1001374	4.43	3	8	2.81	4.93	7.96	7.95	4.24	5.07	5.33	*	+	
MAMMA1001380	1.75	0.68	1.07	1.62	1.08	1.61	1.59	2.67	1.12				
MAMMA1001383	7.51	3.83	5.22	4.97	9.33	8.85	4.6	3.72	6.27				
MAMMA1001394	9.53	6.17	5.52	8.14	10.72	8.86	5.52	7.53	9.35				
MAMMA1001386	24.45	16.7	10.97	12.09	12.45	13	5.97	6.39	7.92				
MAMMA1001389	8.47	4.9	3.19	5.53	5.66	7.55	4.68	4.32	5.33				
MAMMA1001392	6.67	3.9	2.86	4.2	5.48	5.22	3.73	4.27	4.41				
MAMMA1001396	7	4.06	4.91	10.25	16.18	16.77	6.43	5.53	5.34	*	+		
MAMMA1001398	4.11	3.91	3.07	8.57	9.18	8.84	4.16	3.98	4.78	**	+		
MAMMA1001395	5.35	2.58	3.48	7.13	5.55	7.22	4.85	4.29	6.18	*	+		
MAMMA1001399	1.7	1.52	0.97	5.04	3.61	5.38	2.09	2.69	2.33	**	+	+	
MAMMA1001310	10.44	4.9	7.15	8.11	11.15	12.2	4.69	5.11	6.27				
MAMMA1001322	2.58	0.43	0.4	1.79	2.43	1.77	1.08	1.1	1.59				
MAMMA1001324	4.35	2.2	1.73	3.98	4.2	3.77	1.97	2.42	3.42				
MAMMA1001330	13.9	7.33	5.29	11.99	11.29	12.32	6.89	5.13	7.87				
MAMMA1001333	10.64	5.27	5.22	12.45	17.04	13.72	7.27	8.49	8.86	*	+		
MAMMA1001334	19.83	12.61	11.33	16.84	18.12	18.43	11.85	9.39	18.95				
MAMMA1001337	6.8	2.68	3.43	4.92	5.69	6.15	4.3	5.31	5.13				
MAMMA1001341	3.94	2.12	2.51	4.82	3.58	4.32	2.93	4.08	4.66				
MAMMA1001343	4.64	4.02	3.93	10.45	11.27	11.13	3.36	5.55	6.66	**	+		
MAMMA1001344	3.2	1.52	0.8	2.99	5.13	4.05	4.81	3.84	5.02		*	+	
MAMMA1001346	3.61	1.95	1.75	2.88	2.78	3.94	2.71	2.77	4.61				
MAMMA1001350	13.98	5.18	5.89	17.88	22.89	19.58	10.5	8.81	9.31	*	+		
MAMMA1001348	6.8	2.8	3.94	7.53	10.07	7.51	5.93	5.82	6.51				
MAMMA1001326	11.03	6.21	4.6	12.53	13.22	12.6	7.14	6.44	7.15	*	+		
MAMMA1001327	8.15	4.45	6.77	11.06	10.6	9.93	5.76	7.2	5.97	*	+		
MAMMA1001401	12.38	7.29	6.74	14.61	13.5	16.44	10.3	14.7	12.59	*	+		
MAMMA1001408	3.01	1.06	1.25	3.39	2.85	2.94	2.29	2.63	3.03				
MAMMA1001411	13.87	6.35	6.18	6.44	8.45	4.19	7.07	7.42	10.12				
MAMMA1001414	8.9	4.02	3.1	8.97	5.29	6.61	6.05	4.52	6.79				
MAMMA1001415	10.6	3.71	5.04	5.41	5.06	7.32	4.77	5.68	6.24				
MAMMA1001418	5.7	2.73	2.09	6.08	5.21	5.62	4.02	2.75	3.87				
MAMMA1001419	4.73	2.65	2.23	4.77	8	8.11	4.53	3.83	4.07	*	+		
MAMMA1001420	3.1	2.15	1.27	3.76	5.4	5.17	2.79	4.4	3.79	*	+		</

50/180

1578

MAMMA1001534	1.04	1	0.48	0.51	0.82	0.82	0.58	0.71	1.3										
MAMMA1001535	4.92	2.88	1.16	1.88	3.67	4.55	1.49	2.38	2.87										
MAMMA1001547	6.61	3.6	2.98	6.07	6.82	8.95	4.29	5.11	5.04										
MAMMA1001551	6.07	3.86	3.57	4.63	5.65	6.3	4.24	3.97	4.08										
MAMMA1001569	3.5	1.48	2.2	2.86	2.79	2.47	2.33	2.98	1.96										
MAMMA1001575	8.12	4.85	4.3	5.13	5.29	4	4.97	4.91	5.14										
MAMMA1001576	20.26	2.19	9.68	8.21	9.38	6.87	9.09	8.98	9.31										
MAMMA1001584	4.62	2.36	1.31	4.08	5.15	3.32	1.53	1.67	4.4										
MAMMA1001586	1.88	3.47	0.76	1.07	3.5	1.99	1.25	2.13	3.7										
MAMMA1001590	12.7	4.74	4.76	9.14	12.67	13.3	5.6	5.77	7.89										
MAMMA1001599	4.33	1.21	1.88	2.45	2.99	4.36	2.56	2.76	2.04										
MAMMA1001600	5.33	1.77	2.89	2.89	5.09	5.36	2.48	3.86	2.92										
MAMMA1001604	7.87	5.11	1.45	3.42	5.42	5.07	3.4	3.25	5.07										
MAMMA1001606	9.46	4.93	4.75	2.09	8.64	10.49	4.91	6.03	6.85										
MAMMA1001609	2.95	1.2	1.3	2.12	3.38	3.64	2.68	2.56	2.15										
MAMMA1001614	4.39	2.53	1.88	2.49	3.22	3.59	2.48	3.41	3.61										
MAMMA1001615	6.67	1.9	1.82	2.35	2.21	3.65	2.11	2.71	3.83										
MAMMA1001619	19.31	10.08	12.63	10.87	10.6	14.3	14.55	8.6	14.72										
MAMMA1001620	8.92	3.44	4.4	6.63	10.03	12.83	4.85	6.18	5.9										
MAMMA1001623	3.58	4.58	2.08	1.56	2.91	2.34	1.28	2.13	2.52										
MAMMA1001626	2.57	1.13	1.2	1.48	2.12	1.89	1.75	2.77	3.1										
MAMMA1001627	2.24	1.39	0.54	2.13	3.22	2.88	2.13	2.52	2.05										
MAMMA1001630	3.02	5.98	2.09	4.38	4.01	5.45	2.54	3.3	3.8										
MAMMA1001633	6.31	4.02	1.66	7.75	9.37	5.34	5.49	3.61	5.08										
MAMMA1001634	8.31	4.18	4.46	11.22	16.21	13.47	7.21	6.09	6.17	*									+
MAMMA1001635	8.83	4.02	2.32	12.04	8.31	8.32	5.06	3.5	2.52										
MAMMA1001649	4.06	1.62	1.65	3.2	3.67	3.34	1.61	2.68	2.21										
MAMMA1001654	7.5	5.7	4.13	5.16	7.53	6.42	3.33	5.51	3.69										
MAMMA1001660	28.42	20.01	15.26	32.3	33.59	28.79	16.52	14.53	17.32										
MAMMA1001663	16.19	8.13	7.37	24.06	22.04	19.25	11.83	9.81	14.91	*									+
MAMMA1001670	6.04	4.74	3.32	6.72	7.02	6.98	4.35	4.11	5.69	*									+
MAMMA1001671	3.01	0.89	1.47	2.72	3.99	2.13	1.77	2.54	1.32										
MAMMA1001679	4.8	3.29	3	3.03	4.77	2.84	4.71	2.51	4.64										
MAMMA1001683	6.21	3.81	4.22	11.62	10.92	14.02	7.47	6.25	5.71	**									+
MAMMA1001684	1.2	1.06	0.86	1.34	1.65	3.46	1.07	2.23	3.61										
MAMMA1001688	27.08	14.53	17.18	23.31	26.84	30.3	37.53	34.87	43.95				*						+
MAMMA1001689	10.7	4.3	2.46	5.83	12.72	6.26	3.96	2.83	5.24										
MAMMA1001692	5.97	3.39	4.03	11.66	13.26	13.23	4.66	4.11	3.69	**									+
MAMMA1001711	7.12	3.2	3.17	7.6	8.99	7.95	4.59	5.62	7.5										
MAMMA1001715	5.07	1.86	2.28	7.77	5.67	4.34	3.14	3.85	3.95										
MAMMA1001730	5.56	2.96	1.92	1.82	2.04	2.43	2.03	3.01	2.56										
MAMMA1001735	17.93	11.2	11.92	16.49	13.17	19.36	14.97	10.91	15.84										
MAMMA1001740	2.62	1.39	2.19	3.94	5.07	3.81	2.69	2.45	2.08	*									+
MAMMA1001743	63.77	35.5	45.41	34.01	34.01	44.1	49.91	22.06	23.3	*									-
MAMMA1001744	1.18	0.45	0.11	1.34	1.3	0.81	0.46	0.4	0.67										
MAMMA1001745	12.45	7.1	4.31	14.99	16.74	16.98	8.77	5.37	11.73	*									+
MAMMA1001751	5.01	2.42	3.03	4.8	5.52	7.04	3.9	3.22	3.1										
MAMMA1001752	15.56	8.33	10.02	13.09	14.3	13.11	10.96	9.67	11.14										
MAMMA1001754	5.78	4.59	3.53	2.08	6.92	8.14	9.82	5.67	8.59	*									+
MAMMA1001757	1.64	0.65	0.62	1.81	1.16	1.05	0.91	2.59	1.38										
MAMMA1001760	15.19	8.82	7.01	15.51	12.28	21.03	9.85	11.53	17.24										
MAMMA1001764	2.52	1.27	1.35	2.11	2.1	2.28	1.29	2.52	2.02										
MAMMA1001767	3.67	2.6	1.45	4.72	4.48	6.08	3.4	1.82	3.79	*									+
MAMMA1001768	3.4	1.25	1.15	4.85	4.7	4.24	2.45	3.01	3.52	*									+
MAMMA1001769	10.2	4.54	6.07	16.86	18.38	16.58	8.55	6.27	9.46	**									+
MAMMA1001771	7.06	9.36	4.23	3.92	5.03	5.23	5.55	6.69	8.65										

1579

1580

MAMMA1001772	6.61	3.09	3.86	5.22	5.33	3.63	5.11	4.68	6.54										
MAMMA1001773	4.17	2.72	2.42	4.48	7.37	5.12	3.01	4.78	4.22										
MAMMA1001783	6.42	4.36	3.89	10.62	14.19	14.31	4.67	8.23	6.82	**	+								
MAMMA1001785	8.22	2.97	5.14	14.68	12.34	15.26	7.67	8.51	8.54	**	+								
MAMMA1001788	2	0.87	0.27	0.81	1.38	1.73	1.53	0.58	0.8										
MAMMA1001790	5.36	3.86	1.92	6.66	16.36	9.58	3.91	3.27	3.37										
MAMMA1001800	3.52	2.19	1.41	1.85	4.05	2.73	1.44	1.76	1.56										
MAMMA1001804	6.25	3.82	2.87	4.53	3.88	4.64	4.42	4.04	3.96										
MAMMA1001806	3.43	3.08	1.93	7.24	8.78	6.25	3.11	4.51	5.23	**	+								
MAMMA1001812	2.22	1.53	1.51	2.28	2.36	2.64	1.38	2.87	1.34										
MAMMA1001815	1.3	0.41	0.62	2.99	1.2	2.47	2.3	2.24	1.48			*	+						
MAMMA1001817	1.37	3.74	1.14	2.04	2.4	3.09	1.01	1.65	1.29										
MAMMA1001818	2.76	5.34	1.53	1.82	5.05	3.5	2.09	2.95	4.34										
MAMMA1001819	5.52	3.47	3.12	6.33	7.32	6.74	3.51	2.89	5.62	*	+								
MAMMA1001820	2.45	1.25	0.82	2.09	2.1	3.98	4.93	5.44	3.89	**	+								
MAMMA1001824	6.23	3.21	3.26	6.85	6.39	6.61	3.99	4.27	4.97										
MAMMA1001832	3.67	1.55	1.58	4.4	5.34	6.5	1.89	2.88	2.54	*	+								
MAMMA1001834	7.21	6.9	2.37	8.79	8	7.74	7.22	5.59	4.27										
MAMMA1001837	8.71	5.61	5.12	7.73	9.45	10.52	4.01	4.19	6.66										
MAMMA1001848	3.49	1.69	1.44	2.63	4.06	4.52	1.91	2.78	1.99										
MAMMA1001850	20.05	8.18	11.43	18.79	13.27	17.94	12.58	9.7	17.74										
MAMMA1001851	6.25	2.81	2.47	7.34	6.62	10.7	4.31	3.59	5.68										
MAMMA1001852	7.89	5.2	4.18	14.68	10.33	12.24	6.74	5.9	7.65	*	+								
MAMMA1001854	8.11	3.75	3.83	5.47	8.12	7.92	4.25	4.74	5.11										
MAMMA1001858	5.29	6.33	3.33	4.8	9.86	6.77	4.43	4.52	4.66										
MAMMA1001864	6.57	3.87	3.53	5.26	5.92	6.2	4.84	4.25	4.74										
MAMMA1001868	7.13	2.35	1.77	6.07	8.46	12.04	4.49	2.72	4.43										
MAMMA1001874	2.56	0.8	0.99	1.13	2.27	2.32	0.71	0.85	1.82										
MAMMA1001878	14.71	6.24	5.55	12.93	17.25	13.98	8.14	7.86	10.4										
MAMMA1001880	8.73	3.97	3.36	7.33	11.41	9.31	6.98	4.88	7.07										
MAMMA1001885	8.89	4.03	4.1	9.41	9.07	9.64	3.45	4.7	8.89										
MAMMA1001890	10.42	4.8	4.27	13.94	12.16	12.45	5.05	4.52	6.53	*	+								
MAMMA1001893	8.64	3.63	4.1	6.16	5.52	7.2	5.63	4.73	6.76										
MAMMA1001901	3.39	1.13	2.13	3.15	3.75	4.39	2.43	2.45	3.16										
MAMMA1001907	12.12	8.44	5.76	15.43	12.7	15.66	5.86	7.16	6.54	*	+								
MAMMA1001908	16.6	10.48	11.12	10.97	16.32	14.93	6.4	9.69	8.54										
MAMMA1001919	1.82	0.17	0.6	0.94	1.34	0.71	1.26	0.88	0.98										
MAMMA1001931	3.36	2.44	1.38	2.23	3.72	3.2	2.14	2.05	2.86										
MAMMA1001937	5.76	3.91	4.17	7.43	4.75	5.56	4.86	3.34	6.3										
MAMMA1001951	9.42	4.25	4.02	11.76	11.79	12.88	6.81	5.98	6.3	*	+								
MAMMA1001956	12.62	6.26	4.43	11.46	11.33	13.51	7.86	7.63	5										
MAMMA1001957	7.69	6.91	2.97	9.44	10.13	11	3.86	6.71	4.82	*	+								
MAMMA1001960	8.09	4.17	5.2	8.83	7.29	10.11	4.77	4.56	4.66										
MAMMA1001963	1.4	0.45	0.94	0.59	0.92	1.24	0.53	1.56	0.54										
MAMMA1001969	14.58	7.72	8.73	21.99	28.29	25.27	9.69	8.49	9.83	**	+								
MAMMA1001970	13.52	3.54	5.52	13.53	17.34	15.88	8.28	8.88	8.45										
MAMMA1001978	1.45	1.06	0.2	0.2	0.85	0.8	1.52	1.12	0.55										
MAMMA1001992	10.84	5.7	4.65	11.47	10.31	11.31	7.27	6.07	8.17										
MAMMA1001994	10	5.97	3.81	5.9	10.24	11.51	7.66	6.84	4.2										
MAMMA1002008	4.32	3.45	1.54	2.22	2.63	3.21	2.43	4.92	3.14										
MAMMA1002009	6.14	4.06	3.61	6.87	8.92	11.78	3.75	4.94	3.85	*	+								
MAMMA1002011	7.71	3.01	4.35	2.88	2.54	4.06	2.17	2.53	2.34										
MAMMA1002022	5.37	5.17	1.74	4.7	10.07	6.42	3.41	3.09	3.25										
MAMMA1002024	16.93	11.72	9.52	17.19	15.15	15.52	13.99	12.95	15.19										
MAMMA1002032	11.99	7.54	5.7	14.48	16.59	16.42	8.97	9.05	9.69	*	+								
MAMMA1002033	7.72	10.65	3.5	9.82	12.49	7.85	5.26	4.36	7.58										

52/180

【0693】

【表222】

1583

1584

MAMMA1002367	5.13	1.96	2.1	4.1	8	6.58	5.59	7.23	7.33		*	+
MAMMA1002368	4.34	3.93	2.18	3.97	3.15	4.33	1.93	3.77	3.06			
MAMMA1002369	3.43	2.77	0.37	2.27	1.57	2.25	1.64	1.13	1.9			
MAMMA1002370	3.17	4.02	1.28	2.38	4.52	4.47	2.52	2.77	2.84			
MAMMA1002371	8	3.86	4.57	6.11	4.23	6.12	4.47	3.55	4.28			
MAMMA1002372	13.94	6.19	6.42	18.8	17.8	21.12	10.21	8.07	15.59	*	+	
MAMMA1002373	6.97	4.11	3.04	6.45	7.32	6.27	5.03	5.25	5.73			
MAMMA1002374	5.17	2.14	2.44	5.18	5.03	6.05	4.2	2.91	4.33			
MAMMA1002375	5.95	2.63	2	5.32	4.87	5.66	3.33	3.41	4.16			
MAMMA1002376	3.71	2.19	2.17	3.02	3.23	3.18	3.21	2.61	2.25			
MAMMA1002377	4.09	3.82	1.96	6.63	7.73	3.7	2.44	2.99	3.59			
MAMMA1002378	24.32	15.32	19.7	26.21	29.99	31.31	20.38	19.58	18.88	*	+	
MAMMA1002379	10.38	6.59	2.86	14.02	13.82	13.05	10.49	8.04	10.98	*	+	
MAMMA1002380	7.11	4.07	0.96	3.66	5.77	5.39	2.87	2.07	3.97			
MAMMA1002381	5.37	4.98	2.41	6.38	13.31	8.87	4.49	3.92	7.76			
MAMMA1002382	8.07	2.35	5.23	7.19	7.92	8.72	5.3	5.48	6.56			
MAMMA1002383	6.31	4.11	5.15	10.22	11.41	12.06	4.9	7.5	6	**	+	
MAMMA1002384	4.15	2.37	1.67	2.9	3.82	5.04	2.2	3.87	3.47			
MAMMA1002385	4.13	2.74	1.9	3.61	6.19	6.87	2.13	3.26	3.02			
MAMMA1002386	7.26	4	2.1	6.05	5.74	3.04	3.25	4.13	4.42			
MAMMA1002387	10.93	3.6	4.03	10.38	8	8.37	5.57	5.29	6.32			
MAMMA1002388	7.73	3.96	3.73	8.81	10.04	9.43	3.71	3.46	7.48	*	+	
MAMMA1002389	6.93	4.17	2.03	4.83	7.45	7.07	4.3	4.21	4.94			
MAMMA1002390	3.84	3.05	2.4	3.45	5.38	4.65	4.23	5.29	5.91			
MAMMA1002391	5.21	4	2.14	4.04	3.97	4.72	2.11	1.72	2.04			
MAMMA1002392	9.22	7.52	2.31	5.85	8.94	7.55	4.37	4.54	4.03			
MAMMA1002393	5.34	3.25	2.3	4.76	5.27	7.77	2.43	4.79	2.85			
MAMMA1002394	5.57	2.35	1.19	3.19	4.03	4.8	2.05	2.5	2.26			
MAMMA1002395	13.77	9.98	8.17	18.6	20.01	21.01	10.51	7.95	8.5	**	+	
MAMMA1002396	4.19	2.61	1.63	3.14	2.98	2.4	3	1.64	2.41			
MAMMA1002397	6.53	2.69	2.54	6.26	7.25	5.96	4.09	4.49	3.12			
MAMMA1002398	3.93	2.21	1.89	3.56	3.61	4.11	4.72	2.96	3.12			
MAMMA1002399	6.65	2.94	3.45	4.37	4.72	4.67	3.85	4.3	4.84			
MAMMA1002400	7.21	3.57	4.06	7.96	12.17	10.93	5.47	3.81	6.44	*	+	
MAMMA1002401	6.65	2.95	5.07	7.2	8.08	10.65	3.09	4.7	4.45			
MAMMA1002402	4	1.78	2.02	5.31	7.82	7.61	2.14	4.39	2.73	*	+	
MAMMA1002403	1.81	2.58	0.88	2.71	5.37	2.61	2.77	1.86	3.22			
MAMMA1002404	7.22	4.09	4.3	4.23	4.19	3.43	8.27	6.12	7.86			
MAMMA1002405	6.65	3.55	1.7	3.98	7.13	4.08	2.98	3.25	3.05			
MAMMA1002406	10.94	5.98	7.24	14.33	18.89	22.98	6.91	9.41	11.76	*	+	
MAMMA1002407	6.9	2.88	1.85	8.11	6.41	8.49	4.7	4.28	4.05			
MAMMA1002408	1.74	0.88	0.89	1.88	3.53	2.38	2.6	2.64	0.96			
MAMMA1002409	4.98	2.45	2.94	3.63	3.94	4.37	3.81	6.25	5			
MAMMA1002410	5.54	2.15	1.5	3.44	3.65	4.97	2.86	3.08	1.74			
MAMMA1002411	12.21	5.64	2.48	9.88	11.9	8.93	6.13	5.59	4.64			
MAMMA1002412	3.93	2.05	1.27	4.37	4.53	3.05	1.96	4.22	3.47			
MAMMA1002413	6.03	2.26	2.41	5.84	9.22	5	5.51	3.52	6.07			
MAMMA1002414	3.76	1.67	1.82	4.3	5.95	5.66	4.02	3.17	2.93	*	+	
MAMMA1002415	8.04	2.9	2.73	4.67	5.92	6.23	3.94	2.95	5.38			
MAMMA1002416	8.11	3.72	2.87	9.52	10.57	9.58	3.29	5.17	4	*	+	
MAMMA1002417	3.79	2.83	2.72	3.64	5.3	4.09	2.21	3.52	3.36			
MAMMA1002418	6.44	2.97	3.54	5.33	7.97	7.41	2.58	4.01	4.02			
MAMMA1002419	19.95	10.05	7.32	23.49	19.29	16.59	13.96	10.08	15.17			
MAMMA1002420	12.83	8.73	5.05	6.23	10.29	8.7	5.37	6.47	8.29			
MAMMA1002421	8.41	6.54	4.81	4.6	7.71	7.29	6.73	4.8	6.72			
MAMMA1002422	7.42	5.06	2.53	4.57	5.16	4.31	6.56	4.89	5.9			

54/180

【0695】

【表224】

1585

1586

MAMMA1002466	7.61	3.8	3.03	7.05	8.64	7.32	9.99	8.37	11.38		*	+
MAMMA1002479	5.61	2.03	2.45	2.62	3.83	4.24	2.19	2.79	3.07			
MAMMA1002473	2.73	2.58	1.69	4.8	5.81	4.75	1.5	3.35	3.99	**	+	
MAMMA1002489	1.82	0.76	1.1	1.61	2.6	1.72	0.67	1.36	1.72			
MAMMA1002485	11.15	6.59	4.25	5.55	8.76	7.85	6.2	6.28	8.64			
MAMMA1002494	6.22	5.16	3	7.41	9.6	7.67	4.89	3.44	6.03	*	+	
MAMMA1002498	5.71	3.03	1.34	3.92	2.98	3.69	2.66	2.39	3.29			
MAMMA1002534	7.17	3.31	2.26	5.6	4.65	6.85	3.63	4.86	5.05			
MAMMA1002539	5.79	3.23	2.55	4.12	8.81	3.19	5.21	4.47	5.09			
MAMMA1002538	4.01	3.96	2.85	3.37	4.2	2.1	2.88	2.7	3.45			
MAMMA1002545	8.19	4.19	5.05	10.66	9.93	10.97	4.47	4.9	6.19	*	+	
MAMMA1002554	4	1.52	1.49	3.57	3.68	3.97	1.82	2.91	1.1			
MAMMA1002556	9.93	4.82	2.86	7.06	11.34	10.05	3.76	5.07	5.23			
MAMMA1002561	10.06	3.9	4.44	12.05	12.4	15.05	9.97	6.01	8.09	*	+	
MAMMA1002565	4.99	4.2	3.26	4.07	7.56	4.55	3.68	2.91	4.58			
MAMMA1002566	4	2.15	0.94	5.93	2.4	2.55	2.16	2.54	3.99			
MAMMA1002571	7.22	3.36	3.15	5.32	6.04	4.33	4.11	4.2	3.94			
MAMMA1002572	11.2	4.78	6.52	15.53	15.17	13.55	7.02	8.07	9.44	*	+	
MAMMA1002576	6.01	1.71	4.22	10.04	10.33	6.3	4	6.04	6.94			
MAMMA1002584	11.01	7.77	8.72	19.33	19.85	20.62	8.27	12.03	12.19	**	+	
MAMMA1002585	7.85	4.99	2.28	4.43	8.97	3.79	4.39	2.67	4.69			
MAMMA1002586	4.6	2.19	2.47	3.71	4.21	5.32	2.84	2.51	4.3			
MAMMA1002589	4.94	2.94	1.69	6.3	6.89	4.51	3.93	3.36	4.69			
MAMMA1002599	10.71	5.82	7.42	10.33	15.26	8.36	9.91	9.3	15.5			
MAMMA1002593	7.21	1.7	2.9	10.38	6.09	7.62	3.83	4.23	4.78			
MAMMA1002597	5.27	4.72	2.89	5.79	7.99	6.52	3.32	4.98	3.89			
MAMMA1002598	28.18	14.66	17.3	23.76	26.47	26.12	9.35	11.37	10.26			
MAMMA1002603	3.82	2.48	2.87	6.45	7.78	6.16	3.06	4.45	5.16	**	+	
MAMMA1002612	18.88	8.49	7.35	14.76	23.79	19.09	13.04	8.06	15.93			
MAMMA1002617	20.5	11.92	10.78	21.62	26.8	21.46	18.22	10.24	15.46			
MAMMA1002618	8.07	5.37	4.36	5.18	5.81	5.01	3.29	4.53	3.87			
MAMMA1002619	2.75	1.98	1.32	3.42	3.69	3.38	3.52	2.56	2.73	*	+	
MAMMA1002632	4.65	2.19	2.57	6.98	7.16	7	3.88	4.47	5.21	**	+	
MAMMA1002633	3.7	4.09	2.66	8.45	8.43	10.17	4.49	5.06	6.96	**	+	
MAMMA1002635	1.31	0.77	1.1	4.74	4.02	3.9	1.84	3.63	1.92	**	+	
MAMMA1002637	0.15	0.77	0.52	0.63	0.61	1.31	0.61	0.89	0.31			
MAMMA1002639	5	1.49	4.04	8.25	13.1	6.87	3.59	5.41	7.41			
MAMMA1002631	3.02	0.94	0.62	3.54	2.28	2	1.53	1.32	2.73			
MAMMA1002633	8.62	2.1	5.7	4.72	6.74	7.92	3.72	4.69	4.2			
MAMMA1002636	3.99	1.19	1.71	4.59	3.63	5.19	2.99	3.81	3.18			
MAMMA1002637	1.74	1.17	1.01	2.51	1.67	1.58	1.65	2.79	2.2			
MAMMA1002646	5.71	2.6	2.44	4.61	4.24	4.68	2.72	3.67	2.73			
MAMMA1002648	9.62	6.84	5.82	8.64	14.71	12.83	6.98	7.83	7.07			
MAMMA1002659	0.72	0.4	0.49	1.46	0.42	1.02	0.69	0.35	0.84			
MAMMA1002652	6.32	1.69	4.33	6.84	5.22	9.05	3.61	3.81	5.06			
MAMMA1002655	6.13	2.3	1.98	3.61	1.81	5.05	3.34	3.19	3.44			
MAMMA1002662	5.15	2.31	2.11	6.95	6.87	5.4	4.01	4.49	5.25	*	+	
MAMMA1002665	11.8	6.1	10.13	10.87	17.41	15.49	7.23	8.06	7.62			
MAMMA1002671	7.41	2.14	3.42	5.62	4.48	5.33	3.61	3.41	3.76			
MAMMA1002673	7.4	3.46	4.23	7.31	8.7	9.27	5.9	6.54	4.84			
MAMMA1002684	9.53	3.22	5.59	4.24	7.51	8.57	6.73	6.88	7.64			
MAMMA1002685	3.8	1.88	0.7	2.75	4.35	3.69	1.82	1.26	1			
MAMMA1002692	7.2	4.36	3.76	8	7.57	6.47	4.09	3.19	4.9			
MAMMA1002693	8.11	3.16	4.22	9.2	3.75	8.99	4.65	5.78	5.13			
MAMMA1002698	5.29	1.74	2.15	6.64	6.43	7.76	3.35	3.54	3.9	*	+	
MAMMA1002699	2.23	0.61	0.97	1.33	2.22	1.52	1.64	1.71	1.92			

55/180

1587

1588

MAMMA1002701	5.66	2.9	4.33	9.27	7.16	8.59	4.61	5.08	5.08	*	+		
MAMMA1002708	7.94	5.73	7.17	9.47	9.6	11.7	5.3	7.78	6.06	*	+		
MAMMA1002711	5.14	1.55	3.02	5.08	5.35	9.25	4.88	5.17	3.67				
MAMMA1002712	8.23	3.4	3.83	5.92	5.37	4.49	4.33	4.63	3.86				
MAMMA1002716	3.03	1.15	1.75	3.45	3.66	6.18	3.63	4.99	6.27		*	+	
MAMMA1002721	5.09	3.43	2.39	8.57	10.12	9.06	4.73	4.05	4.78	**	+		
MAMMA1002723	3.9	1.75	1.64	3.74	4.55	4.64	2.71	2.75	3.13				
MAMMA1002727	1.94	0.37	0.28	1.65	1.68	1.6	1.31	1.6	1.09				
MAMMA1002728	18.85	12.15	13.58	19.57	15.85	19.98	10.65	11.63	8.96				
MAMMA1002742	24.64	11.73	11.42	17.86	18.78	18.95	12.46	17.75	16.28				
MAMMA1002743	3.32	1.38	1.48	2.64	3.77	2.84	1.3	3.55	2.08				
MAMMA1002744	5	2.18	1.83	8.37	6.2	7.98	3.63	3.32	2.37	*	+		
MAMMA1002746	2.51	0.63	0.79	1.49	2.16	1.83	2.14	1.51	0.81				
MAMMA1002748	3.99	1.96	1.48	3.96	2.53	5.35	2.11	2.64	2.6				
MAMMA1002754	3.27	1.38	1.23	3.72	4.67	3.51	3.5	2.37	3.36				
MAMMA1002758	1.75	1.23	0.68	1.23	1.77	1.88	1.75	1.78	0.81				
MAMMA1002762	15.53	11.07	16.89	14.23	17.23	16.31	8.35	12.66	9.99				
MAMMA1002764	6.2	2.6	2.93	8.75	9.77	8.81	4.73	4.74	4.79	*	+		
MAMMA1002765	4.28	1.57	1.43	2.94	4.93	4.38	2.62	3.87	2.62				
MAMMA1002769	1.56	0.46	0.63	2.76	2.64	1.76	3.07	2.6	2.53	*	+	**	+
MAMMA1002771	7.14	1.91	2.56	3.71	2.39	3.56	2.38	4.39	2.84				
MAMMA1002775	8.17	3.51	3.32	3.63	6.17	5.65	3.96	3.51	3				
MAMMA1002780	4.25	0.67	1.1	3.25	4.36	3.86	1.61	2.45	1.84				
MAMMA1002782	3.73	1.77	1.35	3.47	4.14	4.44	2.59	3.58	3.12				
MAMMA1002795	1.54	0.63	0.41	1.27	1.55	2.07	1.2	2.31	1.82				
MAMMA1002796	5.26	2.04	2.88	2.31	3.68	4.71	3.08	4.01	2.78				
MAMMA1002805	1.95	1.42	2.03	2.66	2.54	2.92	1.33	2.31	1.29	*	+		
MAMMA1002806	7.18	3.13	2.76	7.9	8.06	6.82	4.84	4.21	4.71				
MAMMA1002807	5.28	1.74	0.98	3.68	4.66	5.86	3.42	3.27	3.02				
MAMMA1002814	3.87	2.51	3.12	7.45	7.16	7.74	4.16	4.93	4.92	**	+	*	+
MAMMA1002817	1.7	0.51	0.6	1.42	1.13	1.4	0.99	1.61	0.6				
MAMMA1002820	1.34	1.92	0.86	2.57	2.4	3.83	1.38	1.74	1.69	*	+		
MAMMA1002830	27.11	10.85	16.25	30.04	35.58	32.67	18.44	20.75	20.74	*	+		
MAMMA1002833	6.78	4.02	4.05	10.31	9.78	13.03	4.43	6.24	5.25	*	+		
MAMMA1002835	3.11	0.73	1.29	2.37	4.3	3.68	1.9	2.74	1.11				
MAMMA1002838	5.08	1.94	1.5	7.62	5.02	5.3	2.99	3.7	3.52				
MAMMA1002843	6.45	2.71	2.75	6.39	9.1	5.17	5.25	5.53	5.55				
MAMMA1002843	4.18	1.22	2.78	4.36	3.92	4.27	2.84	3.41	2.54				
MAMMA1002844	15.29	8.97	10.98	13.02	14.25	13.61	12.26	13.86	18.37				
MAMMA1002845	0.94	0.26	0.38	2.62	1.75	2.18	15.33	12.73	11.67	**	+	**	+
MAMMA1002857	92.97	61.45	71.01	93.18	91.48	102.4	49.65	49.13	49.57				
MAMMA1002858	270.3	178.2	193.7	198.5	285	325.3	136.6	154.1	144.4				
MAMMA1002863	6.79	3.17	3.17	4.69	5.56	4.89	3.85	6.3	4.27				
MAMMA1002868	5.34	2.46	2.35	7.72	6.47	7.85	3.3	3.69	4.4	*	+		
MAMMA1002869	6.13	2.1	3.45	4.16	4.01	5.84	3.15	3.68	3.35				
MAMMA1002871	0.97	0.66	0.13	2.7	2.82	2.55	1.36	2.18	2.35	**	+	*	+
MAMMA1002875	4.77	2.06	2.53	6.78	7.19	6.9	3.55	3.8	4.77	*	+		
MAMMA1002879	3.84	2.9	2.39	3.98	5.13	4.33	4.2	4.17	4.62	*	+		
MAMMA1002880	3.28	1.24	0.99	2.01	1.85	1.9	3.12	3.06	1.4				
MAMMA1002881	5.17	2.92	2.09	6.15	9.22	4.65	3.67	4.57	4.68				
MAMMA1002885	5.25	2.83	2.52	4.49	4.87	6.69	2.72	4.39	2.98				
MAMMA1002886	6.24	3.43	2.66	5.52	4.58	6.71	3.64	3.49	2.72				
MAMMA1002887	3.89	0.95	1	1.97	1.75	1.77	2.34	1.93	1.82				
MAMMA1002890	5.13	2.67	3.05	8.31	4.7	8.72	5.27	5.4	5.01				
MAMMA1002892	5.88	3.48	2.47	7.32	8.24	6.42	4.17	5.23	4.57	*	+		
MAMMA1002893	8.86	9.67	8.59	8.18	9.34	9.39	5.69	3.91	5.29	**	-		

56/180

【0697】

【表226】

1590

MAMMA1802895	1.52	1.02	0.66	3.67	2.82	2.63	1.68	3.27	1.67	**	+						
MAMMA1802898	5.3	1.67	2.43	5.04	3.66	3.54	3.19	4.2	4.28								
MAMMA1802905	7.3	4.24	4.9	4.36	3.31	5.5	4.49	4.07	7.6								
MAMMA1802906	7.09	3.55	2.11	4.13	4.15	4.17	3.6	4.08	4.37								
MAMMA1802908	5.1	3.63	2.55	7.12	10.01	7.24	3.97	3.94	6.08*		+						
MAMMA1802909	11.19	2.36	4.9	18.65	20.55	19.49	11.96	9.14	7.19**		+						
MAMMA1802918	8.8	4.28	4.36	7.71	4.97	6.64	4.29	3.85	3.85								
MAMMA1802925	3.35	2.63	1.48	9.46	7.99	8.84	13.12	8.45	14.83*		+	**					
MAMMA1802926	7.82	4.53	3.55	10.54	8.94	10.54	4.02	3.98	2.94*		+						
MAMMA1802939	4.28	1.73	3.17	5.74	5.95	7.07	4.01	5.04	2.57*								
MAMMA1802937	5.96	2.45	3.44	4.74	4.53	7.33	3.19	3.43	4.76								
MAMMA1802938	3.7	2.19	0.47	2.71	4.56	4.15	4.37	4.59	4.01								
MAMMA1802941	1.15	1.12	0.39	3.44	2.75	4.14	1.85	1.74	2.91*		+	*					
MAMMA1802947	6.2	1.75	2	3.69	4.63	4.41	4	2.74	2.53								
MAMMA1802954	3.13	0.8	1.6	3.89	5.54	5.2	2.56	3.32	2.95*		+						
MAMMA1802967	2.77	0.81	0.72	2.65	3.25	3.1	2.15	2.36	2								
MAMMA1802970	10.68	5	6.77	15.62	18.38	19.77	9.12	10.6	10.22**		+						
MAMMA1802971	5.36	1.91	2.72	5.34	4.3	4.54	3.53	5.4	3.71								
MAMMA1802972	3.58	1.23	1.8	5.51	3.48	3.8	2.78	4.51	3.78								
MAMMA1802973	3.05	2.45	2.19	5.84	7.86	5.49	3.04	3.4	3.84*		+						
MAMMA1802979	49.45	21.28	20.21	54.78	50.04	57.56	26.52	29.51	38.14								
MAMMA1802983	1.17	0.84	0.21	1.07	1.04	1.44	0.75	0.85	2.52								
MAMMA1802987	2.51	2.1	1.94	4.65	4.24	4.32	2.66	3.22	2.69**		+						
MAMMA1803003	6.44	2.24	3.39	6.63	8.14	8.81	3.38	3.94	4.55								
MAMMA1803004	2.44	1.12	1.78	4.34	4.64	5.27	2.45	2.33	3.36**		+						
MAMMA1803007	3	0.97	0.37	1.72	3.13	2.66	1.67	2.02	2.34								
MAMMA1803011	6.89	3.86	2.58	10.11	6.23	6.02	5.56	4.68	6.87								
MAMMA1803013	4.71	2.5	3.6	5.96	2.57	4.98	4.47	2.47	4.04								
MAMMA1803015	3.11	1.7	0.83	3.85	3.23	4.39	2.92	3.35	3.6								
MAMMA1803019	1.94	0.48	0.77	1.44	1.99	1	1.47	1.37	1.39								
MAMMA1803020	4.98	3.11	2.83	4.85	4.06	4.94	3.36	4.67	2.34								

57/180

【表 2 2 7】

1591

1592

MAMMA1003140	1.69	0.85	0.6	1.74	1.79	2.62	1.07	1.84	1.32										
MAMMA1003146	3.17	0.64	1.39	1.41	2.11	2.74	2.14	2.47	2.39										
MAMMA1003150	14.6	5.76	6.23	12.63	11.24	8.45	5.72	10.32	7.8										
MAMMA1003154	8.12	5.17	3.61	5.99	6.38	3.91	3.93	4.78	3.8										
MAMMA1003155	3.73	2.43	2.74	2.68	3.2	4.47	4.25	3.56	2.96										
MAMMA1003157	3.72	2.17	1.5	8.43	9.53	5.52	5.81	5.42	4.11	*									
MAMMA1003163	3.24	2.63	2.53	2.86	3.42	4.51	2.32	3.21	3.84										
MAMMA1003164	4.04	1.63	1.78	2.36	3.89	3.12	1.98	3.3	1.9										
MAMMA1003166	2.64	0.97	1.34	1.14	2.03	2.6	0.94	1.46	0.67										
NB9N31000010	14.76	5.71	8.03	2.59	3.65	2.88	2.38	3.3	2.11										
NB9N31000016	7.03	5.06	4.31	4.14	3.19	3.67	2.48	3.3	3.54										
NB9N31000043	6.43	3.37	2.66	3.6	4.63	3.3	4.03	4.8	3.85										
NB9N31000045	19.15	14.02	9.92	7.25	11.2	10.47	9.72	9.74	10.85										
NB9N31000054	6.46	2.26	2.68	6.57	6.74	6.4	6.39	5.01	4.4										
NB9N31000076	2.64	1.86	1.23	4.27	5.28	5.06	3.51	3.29	3.06	**									
NB9N31000086	3.3	1.41	1.24	4.91	5.73	5.78	4.03	3.25	4.37	**									
NT2RM1000001	3.65	2.34	1.78	2.42	3.06	4.27	1.46	2.56	2.65										
NT2RM1000018	18.02	4.88	9.18	11.8	18.97	15.96	10.32	8.58	7.34										
NT2RM1000032	2.53	0.99	1.56	3.18	2.12	2.58	1.32	2.6	2.8										
NT2RM1000035	11.4	5.02	6.42	9.17	9.42	10.51	8.5	7.07	7.86										
NT2RM1000037	13.15	8.99	9.27	10.68	10.22	12.08	8.43	7.97	9.91										
NT2RM1000039	11.18	9.88	11.7	14.16	13.27	16.95	11.97	10.55	15.86	*									
NT2RM1000042	80.13	61.43	48.95	80.07	94.16	101.1	24.69	35.38	37.43										
NT2RM1000055	1.63	0.44	0.19	1.9	1.2	1.06	0.56	1.65	0.56										
NT2RM1000059	10.72	6.4	6.93	10.31	13.85	13	8.96	10.38	10.03										
NT2RM1000063	2	0.27	0.62	1.05	1.09	1.16	1.09	1.18	1.04										
NT2RM1000065	113.3	91.26	69.94	64.48	58.5	52.5	34.11	33.99	50.67										
NT2RM1000066	35.22	18.22	21.68	21.61	23.29	23.48	22.94	24.27	17.75										
NT2RM1000071	63.91	66.46	45.7	62.4	99.26	85.6	34.56	28.87	36.37										
NT2RM1000080	3.9	1.47	1.12	2.18	2.14	2.55	1.54	2.09	2.44										
NT2RM1000086	19.75	10.02	12.84	15.85	21.11	21.57	16.5	12.7	16.82										
NT2RM1000092	3.84	1.47	1.22	4.35	3.45	3.58	5.38	4.65	2.8										
NT2RM1000118	0.16	0.1	0.44	0.44	0.48	0.43	0.45	1.71	0.2										
NT2RM1000119	1.47	0.16	1.14	1.49	1.8	1.77	0.45	3.87	1.63										
NT2RM1000121	3.95	2.18	1.02	2.75	2.63	2.42	2.12	2.47	2.71										
NT2RM1000123	20.69	10.42	10.67	11.66	9.11	15.06	12.71	8.89	10.81										
NT2RM1000127	3.09	0.8	1.57	1.53	1.35	2.79	1.74	2.29	1.61										
NT2RM1000131	1.39	0.57	0.54	0.93	0.82	1.7	1.32	1.99	1.76										
NT2RM1000132	3.41	2.17	2.19	3.36	2.6	3.36	3.07	3.21	1.8										
NT2RM1000153	2.4	1.2	1	2.3	1.9	1.72	2.33	2.75	1.99										
NT2RM1000184	12.46	9.34	11.07	12.61	11.31	13.35	27.02	24.07	25.86	**									
NT2RM1000186	0.96	0.05	1.17	1.92	0.66	0.6	1.01	1.84	0.71										
NT2RM1000187	7.97	7.07	3.88	7.69	10.3	6.3	4.37	5.12	5.93										
NT2RM1000199	2.43	1.17	0.94	2.23	1.56	2.06	2.22	2.21	0.97										
NT2RM1000213	4.77	2.05	1.72	5.31	3.68	5.55	3.01	2.88	2.04										
NT2RM1000215	22.27	12.67	13.12	16.61	13.19	18.25	21.54	17.2	19.32										
NT2RM1000218	4.96	1.49	2.25	6.26	5.56	5.79	6.91	7.15	6.31										
NT2RM1000234	14.47	7.85	6.71	14.79	9.45	14.6	5.58	5.7	6.61										
NT2RM1000236	11.3	7.18	4.01	4.18	3.11	5.39	11.74	17.39	13.83										
NT2RM1000242	-0.07	0.1	-0.14	0.21	0.11	0.96	0.08	1.21	-0.15										
NT2RM1000244	3.77	1.77	0.73	1.27	1.58	1.27	0.95	0.89	0.64										
NT2RM1000252	31.79	17.18	15.85	26.05	28.99	28.97	19.44	15.24	17.84										
NT2RM1000256	20.24	12.26	8.91	9.14	16.96	11.49	13.69	10.22	13.24										
NT2RM1000257	16.34	9.13	9.74	4.83	6.53	7.09	5.1	3.9	3.96	*									
NT2RM1000260	32.33	14.76	16.72	33.82	31.04	33.33	23.69	23.35	23.61										
NT2RM1000269	12.22	8.71	9.44	5.25	3.16	4.72	1.74	2.23	1.16	**									

58/180

【0699】

【表228】

NTZRM1000271	0.75	0.2	0.04	1.21	0.35	0.58	0.94	0.84	0.49										
NTZRM1000272	54.56	36.55	40.59	39.42	48.05	51.89	35.16	41.56	36.18										
NTZRM1000273	25.51	11.38	15.12	14.18	12.87	14.49	8.99	9.27	12										
NTZRM1000274	58.21	39.03	46.94	45.24	44.74	49.05	21.9	22.39	26.39									*	-
NTZRM1000280	3.79	2.05	1.14	3.65	3.57	2.6	4.36	3.9	4.03										
NTZRM1000295	1.04	0.33	0.49	1.43	1.42	1.12	1.49	1.59	1.89	*								*	+
NTZRM1000300	3.37	1.19	1.93	2.35	3.27	3.66	2.84	2	3										
NTZRM1000304	119.7	75.04	105.1	129.6	102.4	124.9	50.36	59.48	58.8									*	-
NTZRM1000314	14.79	10.41	9.09	12.21	10.45	12.98	11.38	9.76	12.93										
NTZRM1000318	24.15	19.1	20.62	18.95	25.93	22.36	13.38	12.74	12.13									**	-
NTZRM1000335	2.7	1.54	1.86	2.64	0.98	2.51	2.11	1.75	0.87										
NTZRM1000341	1.86	1.47	0.19	1.35	0.97	1.03	1.64	1.09	1.69										
NTZRM1000359	12.53	6.61	5.41	9.68	8.63	6.11	10.39	8.69	12.6										
NTZRM1000364	1.42	1.08	1.09	1.11	0.94	2.05	1.14	0.93	0.85										
NTZRM1000355	24.12	12.19	10.53	22.94	22.89	22.53	40.93	26.81	41.82									*	+
NTZRM1000361	3.67	1.47	2.35	2.53	2.08	2.7	1.88	1.68	2.1										
NTZRM1000362	1.06	0.28	0.15	0.8	0.83	1.19	0.3	0.84	1.1										
NTZRM1000372	20.32	11.77	14.09	12.5	15.42	19.07	11.35	13.11	12.12										
NTZRM1000377	4.71	2.13	0.97	3.33	3.33	3.84	3.13	2.47	2.45										
NTZRM1000388	4.08	1.38	1.89	2.94	1.24	2.04	2.06	1.76	3.15										
NTZRM1000394	1.97	0.69	0.13	1.46	1.54	2.03	0.91	0.83	1.86										
NTZRM1000399	1.06	0.34	0.04	1.59	1.17	1.07	1.01	1.52	1.09										
NTZRM1000407	3.28	1.69	1.8	2.92	2.58	2.42	3.74	2.39	2.69										
NTZRM1000421	1.21	0.17	0.31	0.84	0.59	1.24	0.64	0.87	1.2										
NTZRM1000422	184.9	121.2	142.5	178.6	203	174.3	67.17	77.47	67.99									*	-
NTZRM1000438	2.25	0.23	1.58	0.73	1.22	1.54	1.8	1.12	1.6										
NTZRM1000442	11.14	6.84	5.58	14.5	17.82	8.39	4.89	8.25	6.36										
NTZRM1000492	5.37	2.3	2.51	3.94	5.62	7.36	4.89	3.83	3.47										
NTZRM1000512	22.47	26.43	20.07	26.5	33.66	27.9	17.58	19.86	18.1										
NTZRM1000519	29.78	19.56	14.02	7.43	11.19	11.73	14.89	14.37	13.43										
NTZRM1000527	18.16	11.14	6.22	5.88	7.16	7	1.98	1.37	2.55									*	-
NTZRM1000539	12.49	8.93	7.21	6.18	6.43	8.69	2.33	4.94	2.74									*	-
NTZRM1000542	5.88	1.72	2.37	3.23	3.3	5.23	2.07	2.93	2.21										
NTZRM1000543	3.65	0.83	1.64	1.16	1.39	3.69	1.46	2.07	1.37										
NTZRM1000545	54.21	28.45	27.23	49.44	36.73	39.14	24.87	25.09	25.78										
NTZRM1000558	5.67	1.77	2.83	4.02	2.67	3.58	2.91	2.6	1.85										
NTZRM1000563	5.22	2.56	1.89	2.43	2.32	3.96	2.78	2.56	3.17										
NTZRM1000566	7.28	3.71	3.24	1.61	1.5	1.27	1.81	1.72	3.16										
NTZRM1000570	26.49	17.4	16.59	16.76	14.37	17.79	32.95	44.77	33.54									*	+
NTZRM1000571	6.81	1.94	3.76	2.38	2.48	3.22	3.14	4.7	3.91										
NTZRM1000574	1.29	0.74	0.74	1.47	2.46	0.57	1.31	2.11	1.66										
NTZRM1000580	1.69	0.26	0.99	1.9	0.77	2.4	1.57	1.93	1.37										
NTZRM1000620	10.67	5.15	5.67	13.49	19.91	14.9	8.69	7.05	7.31	*									
NTZRM1000623	1.16	0.68	-0.02	1.17	0.81	0.94	0.79	1.05	0.97										
NTZRM1000630	2.05	1.24	0.77	1.67	2.19	1.83	1.87	1.47	1.67										
NTZRM1000633	27.41	17.8	20.59	31.5	32.02	35.16	25.03	10.45	15.13	*									
NTZRM1000634	2.52	1	0.44	1.48	1.48	2.34	1.07	2.17	0.81										
NTZRM1000642	6.47	2.65	3.78	2.95	2.59	5.44	3.92	6.2	4.71										
NTZRM1000647	37.58	16.74	23.8	20.92	20.8	27.77	21.56	20.3	24.27										
NTZRM1000648	2.04	0.41	0.84	1.58	1.79	2.24	1.08	2.71	1.07										
NTZRM1000650	3.85	1.26	1.26	3.06	2.28	2.37	2.52	2.44	1.64										
NTZRM1000661	6.73	4.13	3.51	3.45	3.05	3.31	3.37	4.05	2.46										
NTZRM1000666	23.38	12.8	15.49	0.79	0.92	0.69	0.89	0.98	0.83	*								*	-
NTZRM1000669	3.69	1.54	2.15	2.54	2.09	3.44	2.17	1.76	1.75										
NTZRM1000672	18.91	9.34	13.85	40.77	46.8	48.09	11.37	11.65	12.71	**								+	
NTZRM1000681	7.08	2.25	3.13	16.21	18.15	17.39	29.47	23.33	30.9	**								+	+

1595

1596

NT2RM1000691	1.49	0.33	0.72	2.19	3.8	4.38	1.16	2.44	1.23	*	+		
NT2RM1000692	9.46	4.02	2.95	1.73	2.75	2.69	1.76	3.25	2.56				
NT2RM1000699	5.92	1.52	1.15	3.89	3.52	2.89	2.18	3.43	1.91				
NT2RM1000702	6.62	2.57	3.45	4.5	3.42	3.78	4.09	2.32	3.62				
NT2RM1000703	17.1	15.01	10.3	10.55	11.96	11.61	8.87	8.94	9.74				
NT2RM1000704	65.68	42.42	42.04	15.75	17.49	15.71	13.71	12.1	16.78	*	-	*	-
NT2RM1000725	2.69	1.28	2.86	8.31	19.48	14.98	22.1	28.8	20.7	*	+	**	+
NT2RM1000726	2.12	1.3	1.96	2.34	2.21	3.46	1.65	2.75	1.67				
NT2RM1000731	5.27	2.15	2.93	3.31	4.19	2.99	4.88	3.29	2.95				
NT2RM1000741	1.93	0.67	1.46	0.89	1.2	1.46	1.17	1.5	1.29				
NT2RM1000743	23.68	12.81	12.51	8.34	8.53	8.89	7.58	8.47	7.71				
NT2RM1000744	6.58	2.57	2.31	5.25	4.4	4.66	2.69	3.48	4.72				
NT2RM1000746	6.6	3.69	2.39	2.21	4.12	4.39	2.87	3.97	3.11				
NT2RM1000747	7.04	3.26	3.4	5.08	4.8	5.81	8.95	8.11	9.87		*	+	
NT2RM1000752	2.53	0.89	1.4	2.34	2.42	2.14	1.42	2.26	1.37				
NT2RM1000767	7.61	2.5	4.43	7.29	7.21	8.59	10.72	8.37	9.7		*	+	
NT2RM1000770	5.9	2.04	3.1	5.61	2.94	6.75	3.14	3.37	3.76				
NT2RM1000772	2.24	0.1	0.45	1.66	1.02	0.57	0.12	1.61	0.68				
NT2RM1000779	21.92	14.11	10.14	21.3	25.71	21.61	29.07	21.85	26.85				
NT2RM1000780	3.49	1.84	0.6	4.74	3.37	4.7	3.33	3.29	1.67				
NT2RM1000781	0.57	0.24	0.41	1.11	0.76	1.25	0.94	2.16	0.86	*	+		
NT2RM1000789	3.24	2.46	2.34	3.02	3.98	4.62	2.09	4.84	3.17				
NT2RM1000800	7.44	3.44	7.11	8.01	9.85	8.74	6.51	5.53	7.87				
NT2RM1000802	9.35	5.25	6.84	5.12	5.47	5.85	9.59	9.36	9.75				
NT2RM1000811	0.9	0.16	0.89	1.36	1.11	1.28	0.91	1.35	0.23				
NT2RM1000826	26.11	13.59	16.15	23.62	25.62	25.75	12.43	12.08	10.27				
NT2RM1000839	4.42	3.56	2.62	8.2	6.8	9.18	6.67	6.07	6.37	**	+	**	+
NT2RM1000851	96.56	76.65	61.3	78.41	75.7	87	48.08	33.56	47.08	*	-		
NT2RM1000853	6.27	2.21	1.64	3.09	3.54	4.73	6.47	7.68	4.1				
NT2RM1000854	4.84	2.51	2.09	5.62	3.9	3.49	3.8	5.68	4.28				
NT2RM1000841	32.04	19.08	20.07	17.66	18.86	19.57	17.83	9.4	13.52				
NT2RM1000848	22.37	12.31	11.25	14.54	11.17	13.09	8.36	10.63	15.1				
NT2RM1000850	1.25	0.36	0.94	1.01	0.67	1.33	1.5	1.94	1.75		*	+	
NT2RM1000852	3.74	0.76	1.24	2.68	2.43	2.34	2.39	3.1	1.87				
NT2RM1000853	1.46	0.57	0.14	1.6	2.87	1.74	1.25	0.52	1.87				
NT2RM1000855	19.04	8.47	10.06	15.32	18.2	15.69	26.5	18.76	20.5				
NT2RM1000857	20.9	10.06	10.76	20.92	27.84	24.62	16.83	13.46	17.36				
NT2RM1000858	22.68	8.04	9.94	22.93	26.24	26.47	20.88	15.02	18.54				
NT2RM1000867	15.69	9.11	9.26	15.56	10.14	14.92	15.07	11.26	10.73				
NT2RM1000874	9.77	5.6	5.03	6.49	6.79	8.79	8.74	7.92	8.94				
NT2RM1000883	4.01	2.76	2.65	5.69	5.23	6.94	2.13	4.39	2.7	*	+		
NT2RM1000883	17.32	10.68	13.68	15.2	15.74	17.32	14.61	9.93	20.96				
NT2RM1000885	31.05	13.08	10.39	19.2	20.71	27.92	20.23	18.36	23.03				
NT2RM1000892	3.73	1.65	2.82	3.47	1.63	2.22	4.97	4.49	6.3		*	+	
NT2RM1000894	14.4	9.62	11.92	7.88	9.3	10.29	9.51	9.36	13.18				
NT2RM1000898	2.53	0.85	1.96	3.01	2.71	4.11	3.76	3.77	6.2		*	+	
NT2RM1000899	1.45	0.26	1.26	1.48	1.2	1.14	1.07	1.69	0.72				
NT2RM1000905	55.04	22.33	30.63	36.24	41.24	41.41	17.87	22.74	23.5				
NT2RM1000920	7.05	2.93	6.34	6.29	7.41	5.83	7.31	6.05	5.79				
NT2RM1000914	8.32	4.94	2.32	6.53	12.83	8.37	4.34	8.86	6.23				
NT2RM1000919	4.63	2.11	2.49	5.45	3.02	4.5	2.58	2.81	4.74				
NT2RM1000921	2.3	0.73	0.47	1.57	1	2.01	1.98	1.39	1.88				
NT2RM1000922	7.7	4.51	3.3	6.07	5.41	6.35	3.4	3.21	3.38				
NT2RM1000924	3.39	1.7	1.15	2.35	2.35	2.87	1.24	2.25	1.77				
NT2RM1000927	3.83	1.15	1.76	5.16	2.77	6.27	2.3	3.24	2.14				
NT2RM1000931	8.45	4.91	4.93	9.07	6.69	6.96	5.29	7.06	3.5				

60/180

【0701】

【表230】

1599

1600

NT2RM2000259	6.85	2.87	3.45	6.74	6.95	8.42	4.64	4.72	5.57										
NT2RM2000259	9.6	4.08	4.77	6.02	9.47	7.13	5.19	6.42	6.9										
NT2RM2000260	9.93	9.2	6.51	4.88	7.9	8.73	11.23	8.04	9.57										
NT2RM2000265	2.4	1.14	0.66	1.28	0.86	1.86	1.3	1.27	1.08										
NT2RM2000287	10.73	4.68	6.12	10.38	10.35	12.59	6.93	10.27	8.06										
NT2RM2000306	16.48	15.91	13.02	16.75	16.33	10.75	17.88	8.38	16.11										
NT2RM2000312	57.19	46.28	42.21	59.66	41.08	60.14	43.74	21.02	32.47										
NT2RM2000322	6.45	2.73	3.3	5.49	4.98	2.77	3.63	4.55	3.78										
NT2RM2000343	5.35	4.3	5.69	10.01	10.47	9.81	6.04	4.74	6.91**	+									
NT2RM2000359	5.94	2.73	3.95	5.3	4.77	4.66	3.08	3.35	2.77										
NT2RM2000362	15.37	16.06	11.14	15.03	19.07	17.41	12.3	11.08	9.04										
NT2RM2000363	2.27	1.12	1.53	3.15	1.57	1.39	1.27	1.95	1.06										
NT2RM2000368	20.14	10.44	9.67	11.84	14.77	11.87	10.3	9.5	10.03										
NT2RM2000371	111	74.6	73.79	116.3	62.15	121.3	50.3	42.75	56.01										
NT2RM2000374	4.78	2.52	1.94	6.65	5.32	5.42	4.66	3.93	3.68*	+									
NT2RM2000387	11.91	6.37	5.79	20.24	13.27	20.63	9.51	12.58	11.14*	+									
NT2RM2000393	3.45	1.01	1.83	2.71	1.61	3.18	1.81	3.53	1.53										
NT2RM2000395	1.44	0.49	0.91	2.24	0.76	1.26	1.08	2.52	0.72										
NT2RM2000402	7.26	1.87	2.95	6.33	6.77	7.71	5.51	6.64	5.38										
NT2RM2000405	5.34	2.42	2.76	3.26	3.78	4.88	2.25	2.56	2.19										
NT2RM2000407	19.34	9.37	10.6	5.59	9.51	9.38	8.65	7.51	10.04										
NT2RM2000410	3.06	1.14	0.97	2.09	2.96	2.28	2.57	1.94	2.16										
NT2RM2000420	4.52	1.56	1.71	6.72	7.81	5.85	4.96	3.72	3.6*	+									
NT2RM2000422	14.32	4.96	7.79	15.68	12.45	9.99	14.38	10.45	10.29										
NT2RM2000423	3.93	2.29	3.18	9.3	10.31	11.58	4.01	3.67	2.37**	+									
NT2RM2000453	4.1	1.67	3.69	10.71	9.43	9.18	6.96	4.45	5.45**	+									
NT2RM2000469	1.22	0.59	0.27	2.22	1.54	1.32	1.52	1.06	1.82										
NT2RM2000480	4.98	2.59	1.93	4.39	4.04	3.10	5.95	3.52	4.92										
NT2RM2000497	2.77	1.77	1.58	7.44	5.74	5.87	2.86	3.26	4.3**	+									
NT2RM2000502	4.18	2.99	2.68	7.32	4.36	3.54	3.69	2.68	5.35										
NT2RM2000504	2.49	1.56	2.01	5.06	3.93	4.92	5.83	4.60	4.88**	+	++								
NT2RM2000514	5.60	3.19	3.45	8.34	7.66	5.47	4.66	4.70	6.69										
NT2RM2000522	0.63	0.58	0.61	1.36	0.80	1.01	0.53	0.67	1.87										
NT2RM2000540	5.03	4.07	2.80	5.25	6.86	2.78	4.31	3.32	4.3										
NT2RM2000556	0.38	0.75	0.50	1.40	1.96	0.69	3.19	0.77	0.73										
NT2RM2000565	4.89	2.53	3.37	4.40	4.50	4.25	5.66	3.06	4.57										
NT2RM2000566	5.85	4.38	3.46	8.37	5.27	4.67	4.65	4.38	5.92										
NT2RM2000567	4.29	3.03	2.89	4.78	3.00	1.68	3.19	2.38	4.64										
NT2RM2000569	6.50	3.15	2.85	8.65	8.54	6.48	4.57	3.91	4.43										
NT2RM2000577	11.83	4.68	6.45	6.50	8.99	3.96	4.84	6.67	8.79										
NT2RM2000581	6.47	3.33	5.21	7.46	8.40	4.99	4.74	5.34	7.76										
NT2RM2000582	5.88	3.81	3.49	9.44	7.98	6.09	7.69	6.61	8.15*	+	*								
NT2RM2000583	22.92	13.30	11.99	23.97	16.17	19.54	16.8	11.46	18.28										
NT2RM2000589	11.18	6.26	6.74	9.54	8.57	7.04	5.39	6.22	7.18										
NT2RM2000594	11.31	9.59	11.31	3.91	4.21	3.25	3.27	3.48	2.37**	-	**								
NT2RM2000599	22.01	15.12	17.66	19.78	24.09	21.66	15.2	13.93	15.24										
NT2RM2000609	2.49	1.70	2.43	4.47	3.94	3.24	1.96	3.24	2.3*	+									
NT2RM2000612	3.82	2.55	2.84	4.46	7.55	3.95	4.78	3.73	4.27										
NT2RM2000622	8.85	7.06	10.37	13.55	16.80	10.42	7.83	8.92	10.48										
NT2RM2000623	23.78	13.14	15.60	19.91	22.53	23.26	22.22	15.23	19.45										
NT2RM2000624	16.48	10.64	4.76	11.37	17.12	10.30	8.76	8.23	137.7										
NT2RM2000632	5.44	2.83	2.35	3.85	3.76	2.79	2.22	2.21	7.42										
NT2RM2000635	2.91	2.32	2.35	7.82	9.57	5.76	5	4.36	4.83**	+	**	+							
NT2RM2000636	3.87	2.82	3.19	5.69	5.77	3.68	4.63	3.86	4.55										
NT2RM2000639	4.56	3.86	3.29	4.47	7.45	4.02	3.6	6.93	4.67										
NT2RM2000649	4.09	4.81	3.74	4.86	8.90	5.64	4.74	3.82	6.57										

62/180

【0703】

【表232】

1601

1602

NT2RM2000658	7.80	7.19	11.39	10.12	9.62	7.80	6.87	5.57	7.23										
NT2RM2000660	27.64	11.87	13.50	20.31	25.06	18.91	13.54	13.65	15.55										
NT2RM2000669	7.79	4.71	4.17	9.97	13.43	8.55	3.67	4.50	6.66										
NT2RM2000689	29.82	30.60	28.82	42.51	72.34	53.67	22.11	19.71	38.62	*	+								
NT2RM2000691	4.67	3.54	3.74	5.23	6.41	4.14	4.29	3.98	4.19										
NT2RM2000714	13.27	8.60	10.19	9.82	10.81	9.42	13.37	9.65	17.53										
NT2RM2000718	1.36	1.54	1.09	3.28	7.10	3.02	2.42	2.48	2.19			**	+						
NT2RM2000732	6.10	4.20	5.69	12.72	15.74	11.49	5.77	6.42	7.79	**	+								
NT2RM2000735	24.38	15.21	20.46	56.19	49.62	47.05	16.37	24.66	27.14	**	+								
NT2RM2000740	6.48	2.95	2.62	6.53	5.49	3.44	3.93	3.46	2.74										
NT2RM2000743	21.35	12.67	14.35	10.73	9.73	9.68	2.24	1.81	2.16			**	-						
NT2RM2000772	11.89	7.81	9.52	17.15	14.77	14.45	6.23	7.95	10.31	*	+								
NT2RM2000773	11.75	6.40	6.69	9.73	11.32	9.29	9.82	8.51	8.01										
NT2RM2000776	12.66	6.48	11.36	17.08	19.56	14.42	12.22	8.19	11.56	*	+								
NT2RM2000784	11.22	7.09	6.83	7.88	10.63	6.42	6.22	6.90	7.64										
NT2RM2000795	9.52	5.29	6.34	17.74	18.61	15.80	6.53	8.43	10.09	**	+								
NT2RM2000796	27.57	17.52	26.46	2.02	2.40	3.17	1.82	2.65	1.66	**	-	**	-						
NT2RM2000798	14.84	8.16	10.91	45.29	27.47	24.14	26.69	20.97	28.82	*	+	**	+						
NT2RM2000801	37.70	23.20	28.38	26.35	37.85	28.51	31.37	32.22	38.5										
NT2RM2000821	3.67	2.04	2.27	8.85	6.90	6.15	5.86	5.63	5.4	**	+	**	+						
NT2RM2000829	36.66	22.85	41.47	29.93	23.94	16.17	15.48	17.92	19.23	*	-								
NT2RM2000837	5.77	3.15	3.99	6.12	6.76	5.46	5.15	4.55	4.39										
NT2RM2000924	6.69	5.13	4.70	12.18	14.72	8.21	5.5	6.80	8.89	*	+								
NT2RM2000930	14.27	7.36	9.58	15.72	15.41	13.15	7.93	7.73	11.49										
NT2RM2000937	2.93	2.09	3.52	5.00	4.64	3.14	1.89	3.58	2.8										
NT2RM2000939	6.56	3.88	4.32	5.94	7.25	6.23	4.34	5.73	5.56										
NT2RM2000943	141.00	79.29	113.17	107.50	122.19	108.41	73.07	66.91	67.18										
NT2RM2000951	4.09	2.69	2.78	3.88	3.40	4.39	3.48	3.83	3.33										
NT2RM2000952	5.14	3.58	3.50	6.02	4.82	4.48	3.55	3.67	3.9										
NT2RM2000966	11.75	10.12	10.87	9.00	11.41	11.06	9.18	10.30	5.82										
NT2RM2000973	22.49	16.16	17.58	24.24	28.57	21.97	14.32	17.17	15.94										
NT2RM2000983	10.51	6.87	10.06	15.15	16.05	11.81	9.62	13.40	12.47	*	+								
NT2RM2000984	3.34	2.49	1.94	4.17	6.33	3.91	3.14	3.09	3.89										
NT2RM2000994	17.72	5.91	15.58	25.00	22.32	16.64	8.13	8.32	6.15										
NT2RM2001004	6.95	4.49	3.43	6.09	8.10	5.86	5.16	4.92	6.83										
NT2RM2001023	113.50	66.21	87.63	148.44	181.02	157.90	78.72	73.28	91.6	*	+								
NT2RM2001035	10.78	6.86	10.47	14.95	15.69	13.90	7.29	8.73	9.42	*	+								
NT2RM2001038	4.09	2.22	2.89	6.55	5.43	6.97	3.62	3.51	3.32	*	+								
NT2RM2001043	2.10	1.71	2.70	4.88	5.53	4.13	3.52	4.59	4.54	**	+	*	+						
NT2RM2001050	8.66	4.61	6.50	7.54	9.45	9.85	5.61	5.16	6.52										
NT2RM2001053	4.62	4.14	3.41	6.16	3.15	5.46	4.13	4.67	4.8	*	+								
NT2RM2001065	6.07	2.63	3.08	8.01	7.85	5.22	3.46	3.40	2.98										
NT2RM2001075	101.53	60.27	56.87	59.75	60.87	48.63	40.45	36.79	33.7										
NT2RM2001083	13.68	8.75	8.30	8.28	9.63	7.55	10.14	7.92	7.18										
NT2RM2001100	8.62	6.13	5.38	7.77	11.80	7.92	8.12	5.53	6.12										
NT2RM2001105	18.36	12.31	11.09	26.95	28.47	25.34	13.8	13.76	12.91	**	+								
NT2RM2001109	5.91	3.28	4.91	5.36	6.02	5.10	4.49	5.78	4.86										
NT2RM2001110	9.13	5.14	5.81	7.93	9.23	8.95	5.5	5.40	7.45										
NT2RM2001126	4.23	4.04	4.69	10.78	10.09	7.28	4.72	4.73	5.55	**	+								
NT2RM2001131	9.35	4.34	5.26	6.74	7.12	6.44	4.76	4.21	3.9										
NT2RM2001141	9.27	7.43	7.38	17.22	17.01	12.50	8.53	7.89	8.62	*	+								
NT2RM2001152	3.64	1.47	1.46	2.09	3.47	2.45	1.42	1.89	2.81										
NT2RM2001177	8.38	4.92	5.00	10.70	11.58	8.68	5.53	6.90	6.04	*	+								
NT2RM2001194	10.76	6.38	8.60	11.33	15.08	9.90	8.38	9.20	9.42										
NT2RM2001195	3.62	3.00	3.18	4.85	7.13	3.67	3.45	3.91	3.64										
NT2RM2001196	7.18	4.57	6.50	10.22	17.76	12.85	4.98	6.12	9.71	*	+								

63/180

【0704】

【表233】

1603

1604

NT2RM2001201	13.08	8.55	9.63	10.72	11.46	9.31	8.37	10.02	9.79										
NT2RM2001231	6.92	2.79	3.15	5.91	7.22	4.72	4.61	5.13	3.98										
NT2RM2001238	2.81	1.05	1.43	3.40	2.72	2.10	1.81	2.65	3.37										
NT2RM2001243	6.98	4.99	5.16	9.29	9.00	6.32	4.34	5.08	4.64										
NT2RM2001244	4.98	5.59	4.41	14.49	19.11	7.34	5.11	6.41	7.87										
NT2RM2001247	15.41	9.79	11.87	12.82	15.98	10.20	6.66	8.32	9.67										
NT2RM2001256	2.93	2.70	3.12	2.39	2.54	2.02	2.24	3.49	2.22*										
NT2RM2001269	1.76	1.73	1.47	3.07	6.49	3.10	1.39	5.05	2.29										
NT2RM2001278	7.64	6.14	6.38	12.27	11.97	10.88	6.39	7.92	7.27**										
NT2RM2001291	4.14	2.35	1.90	4.62	4.03	2.79	3.65	2.48	3.16										
NT2RM2001294	10.67	6.20	5.16	12.58	9.68	9.06	8.36	5.49	6.33										
NT2RM2001295	4.70	3.78	3.23	5.43	4.66	4.21	4.46	4.14	4.92										
NT2RM2001302	5.63	4.69	4.19	1.74	2.61	0.97	2.97	3.64	4.24**										
NT2RM2001306	2.52	1.56	1.39	3.47	5.32	4.74	2.64	2.44	2.72*										
NT2RM2001312	1.22	1.12	0.35	2.84	2.71	1.41	1.09	2.09	1.77										
NT2RM2001319	5.09	3.21	4.08	5.71	5.46	5.01	3.84	5.43	5.66										
NT2RM2001324	8.85	3.42	3.83	7.05	8.29	8.06	5.36	6.31	4.89										
NT2RM2001345	12.36	6.03	4.96	4.58	10.06	7.26	10.14	5.50	8.06										
NT2RM2001360	9.69	4.48	4.35	8.36	5.80	5.82	6.45	4.63	6.16										
NT2RM2001379	1.53	1.04	0.81	1.70	1.86	1.18	1.6	2.44	2.47										
NT2RM2001391	1.02	1.38	1.05	3.81	3.30	1.71	1.72	1.73	1.75*										
NT2RM2001393	6.61	4.78	7.01	5.53	6.68	4.32	4.86	4.39	4.92										
NT2RM2001420	2.35	0.95	1.41	3.00	4.15	1.59	1.98	2.45	1.71										
NT2RM2001423	11.93	5.27	6.94	5.99	7.80	3.34	2.15	4.14	4.71										
NT2RM2001424	18.20	9.15	9.42	11.35	10.96	8.30	11.11	9.35	12.67										
NT2RM2001482	15.21	7.55	7.78	14.57	12.13	9.92	11.31	8.31	11.15										
NT2RM2001499	16.92	9.02	7.05	8.26	6.45	6.32	5.19	4.43	7.42										
NT2RM2001504	3.91	2.51	1.97	4.23	4.34	3.86	4.03	2.84	4.42										
NT2RM2001524	2.28	1.47	1.87	2.95	3.08	2.80	2.63	3.34	2.29*										
NT2RM2001530	0.78	0.43	0.54	2.16	2.44	1.43	1.65	1.93	1.93*										
NT2RM2001533	5.77	3.13	3.08	6.59	7.98	5.62	5.57	5.84	5.16										
NT2RM2001540	29.91	19.29	20.03	25.11	24.66	12.51	8.93	9.56	11.82										
NT2RM2001544	5.22	2.70	2.16	5.77	5.72	5.99	4.13	3.93	3.57										
NT2RM2001547	10.18	3.47	3.29	5.82	9.93	4.61	8.42	7.52	11.22										
NT2RM2001558	4.96	2.25	2.36	3.07	3.85	4.04	4.67	2.71	4.49										
NT2RM2001575	4.76	2.31	3.04	7.85	7.43	4.47	3.66	3.23	5.49										
NT2RM2001583	3.25	3.39	2.40	5.62	5.69	4.66	5.53	3.88	4.63**										
NT2RM2001588	2.97	1.41	1.47	4.20	4.38	3.50	3.05	3.37	3.85*										
NT2RM2001592	1.95	2.06	1.67	3.66	3.58	2.66	2.98	2.38	2.72*										
NT2RM2001603	7.68	4.12	5.43	8.07	9.92	5.79	4.3	6.43	7.62										
NT2RM2001605	6.36	3.57	2.87	8.10	9.32	7.63	6.11	4.82	7.04*										
NT2RM2001611	4.43	2.58	2.01	5.92	8.58	4.85	5.15	3.69	4.23										
NT2RM2001613	5.87	2.94	3.70	6.48	9.87	7.29	8.01	9.10	11.64										
NT2RM2001624	11.27	3.06	6.34	7.63	9.90	5.32	10.52	8.42	10.76										
NT2RM2001632	8.60	4.62	8.41	12.48	14.32	11.43	11.18	11.01	12.96*										
NT2RM2001633	1.62	1.36	1.29	4.23	3.15	3.36	2.92	3.34	2.97**										
NT2RM2001635	6.76	5.69	4.97	6.78	9.41	8.26	7.11	8.37	8.13										
NT2RM2001636	4.43	3.06	3.83	4.52	5.10	3.16	3.81	3.42	3.49										
NT2RM2001637	2.79	1.78	2.31	4.20	4.67	2.96	3.03	3.93	2.28*										
NT2RM2001639	4.58	2.63	2.19	3.05	3.54	3.49	2.88	2.16	3.34										
NT2RM2001641	3.30	2.69	1.81	2.72	3.01	1.92	1.93	2.95	2.91										
NT2RM2001643	3.00	1.41	2.34	4.92	3.73	2.89	3.15	3.35	3.43										
NT2RM2001648	3.60	1.94	2.50	5.96	6.92	4.37	6.52	6.11	8.21*										
NT2RM2001652	4.13	2.45	1.80	4.68	5.72	3.86	2.23	2.96	4.29										
NT2RM2001659	1.81	1.41	1.26	2.31	1.88	1.34	2.02	2.75	2.32										
NT2RM2001660	2.12	1.41	1.99	2.87	4.60	2.23	2.13	3.03	3.35										

64/180

【0705】

【表234】

1606

NT2RM2801664	5.67	1.74	2.53	4.44	5.25	4.88	2.57	2.77	3.63								
NT2RM2801668	7.83	4.11	5.80	12.91	11.03	10.32	6.9	5.64	6.89	*	+						
NT2RM2801670	5.07	2.93	3.77	4.21	4.81	3.14	3.67	3.52	4.76								
NT2RM2801671	2.26	2.13	2.75	6.03	4.05	5.74	4.08	5.29	5.9	*	+	**	+				
NT2RM2801675	0.53	0.71	0.81	1.96	1.15	1.66	0.84	1.96	0.76	*	+						
NT2RM2801681	1.11	1.22	1.01	3.34	4.29	2.27	1.69	3.16	1.72	*	+						
NT2RM2801685	3.03	2.26	1.29	2.06	2.47	1.90	1.92	3.02	2.65								
NT2RM2801689	2.78	1.66	2.54	4.45	4.23	2.30	3.29	3.27	2.72								
NT2RM2801695	7.30	3.32	3.64	20.95	20.35	18.16	12.07	10.51	12.36	**	+	**	+				
NT2RM2801696	13.28	6.12	3.86	8.81	10.82	9.78	6.65	6.44	6.65								
NT2RM2801699	8.16	3.47	3.88	5.88	6.34	6.37	6.66	7.84	5.32								
NT2RM2801699	2.40	2.32	1.42	3.33	3.59	3.21	1.64	3.47	3.24	*	+						
NT2RM2801700	2.41	1.38	1.00	2.93	2.03	1.36	1.5	2.70	2.35								
NT2RM2801704	6.94	4.34	5.63	17.99	22.84	16.16	12.13	12.06	13.82	**	+	**	+				
NT2RM2801706	1.19	2.60	4.07	6.98	8.64	6.26	3.29	5.55	5.04	*	+						
NT2RM2801714	1.72	1.75	2.15	3.05	3.64	2.23	1.71	2.86	4.84								
NT2RM2801716	16.89	6.66	8.99	10.52	14.37	8.33	10.03	10.20	10.62								
NT2RM2801718	13.66	7.01	6.41	14.04	11.83	11.25	5.12	7.15	10.53								
NT2RM2801723	6.13	3.06	3.78	9.65	9.89	7.73	4.12	5.35	3.42	*	+						
NT2RM2801727	5.93	4.01	4.52	4.87	5.62	3.99	5.43	6.14	6.96								
NT2RM2801730	3.02	1.57	1.66	3.08	3.68	4.44	2.79	3.79	2.6								
NT2RM2801738	6.78	3.40	5.60	4.93	5.41	3.52	4.55	4.50	4.55								
NT2RM2801743	4.12	2.65	1.97	3.64	5.10	2.62	3.21	3.25	2.82								
NT2RM2801753	4.87	2.89	3.87	7.06	7.68	7.46	4.96	5.77	5.73	**	+						
NT2RM2801755	11.15	5.43	7.63	8.83	12.50	9.88	7.94	7.72	5.34								
NT2RM2801760	6.52	3.36	4.22	8.42	9.37	6.40	10.28	9.84	11.76								
NT2RM2801765	1.13	1.98	1.79	3.23	3.97	3.48	2.65	2.41	2.82	**	+	*	+				
NT2RM2801767	12.87	8.82	9.72	11.08	15.03	8.12	9.19	9.22	14.64								
NT2RM2801768	3.41	2.58	3.68	3.47	6.28	4.04	2.49	2.74	3.01								
NT2RM2801771	4.11	3.62	4.50	11.05	10.86	9.39	5.06	5.82	8.71	**	+						
NT2RM2801778	1.70	1.61	1.19	3.14	4.69	2.67	2.01	2.74	1.97	*	+						
NT2RM2801782	3.27	3.78	3.39	3.01	5.59	4.13	3.83	4.97	5.07					*	+		
NT2RM2801784	3.64	1.97	1.45	2.55	4.38	1.85	2.15	2.16	2.26								
NT2RM2801785	11.40	5.25	4.67	8.49	7.03	6.72	4.99	4.72	4.92								
NT2RM2801792	5.79	3.39	4.17	6.99	5.40	4.24	3.59	5.22	5.39								
NT2RM2801795	9.85	4.56	3.32	7.91	9.48	5.77	7.27	6.25	5.93								
NT2RM2801797	5.04	2.64	2.13	7.82	15.93	10.34	3.54	4.95	3.54	*	+						
NT2RM2801800	3.26	2.51	2.46	4.20	4.38	3.21	2.99	3.72	2.42								
NT2RM2801803	3.60	2.31	2.65	4.14	6.89	5.00	2.04	3.10	3.17								
NT2RM2801905	1.03	0.92	2.17	2.21	3.99	1.67	0.87	3.16	1.79								
NT2RM2801906	5.77	1.94	1.66	4.46	3.73	2.85	3.42	3.44	3.44								
NT2RM2801913	3.38	1.75	1.74	2.55	3.99	2.42	1.83	1.59	3.71								
NT2RM2801914	3.09	1.71	2.83	3.06	4.28	2.96	1.96	3.02	3.47								
NT2RM2801918	2.38	1.33	1.54	3.40	2.50	2.32	1.89	3.32	1.89								
NT2RM2801923	1.26	1.12	0.39	0.95	1.88	0.91	0.96	2.06	1.08								
NT2RM2801925	10.44	6.78	7.52	10.86	11.22	8.43	10.34	16.17	16.27			*	+				
NT2RM2801933	4.52	2.18	1.93	4.11	5.31	3.71	1.98	4.92	3.68								
NT2RM2801939	16.50	9.01	12.64	20.38	24.01	12.42	26.45	40.89	49.99			*	+				
NT2RM2801940	7.75	3.07	2.83	17.33	13.18	10.65	6.84	6.84	7.97	*	+						
NT2RM2801951	7.34	4.30	5.43	12.87	13.61	10.39	7.97	4.60	6.49	**	+						
NT2RM2801955	5.55	3.48	2.68	4.96	5.56	4.20	7.12	5.33	6.29								
NT2RM2801967	3.35	3.93	2.06	3.68	4.49	3.19	3.59	2.29	4.09								
NT2RM2801969	28.84	23.52	26.51	34.13	35.79	25.24	15.38	12.75	18.43			**	-				
NT2RM2801979	0.65	1.04	0.48	2.26	1.68	1.53	1.33	1.12	1.72	*	+	*	+				
NT2RM2801983	3.25	3.47	2.90	9.89	10.82	12.52	5.43	4.02	7.24	*	+						
NT2RM2801986	2.86	1.25	2.11	3.84	5.06	2.09	1.79	1.91	1.93								

65/180

1608

NTZRM2001887	4.05	2.53	2.07	3.94	3.93	2.72	2.74	2.00	2.87										
NTZRM2001896	968.51	557.14	625.69	446.49	419.99	290.65	817.5	613.90	955.7										
NTZRM2001902	1.32	1.90	1.03	2.63	3.33	2.08	2.5	1.84	1.37	*	+								
NTZRM2001903	10.52	8.17	6.65	10.52	9.78	8.75	7	6.78	10.05										
NTZRM2001909	5.61	3.44	3.21	5.48	6.96	3.46	4.44	4.85	6										
NTZRM2001935	3.82	1.91	1.54	3.50	4.79	3.97	2.7	3.73	4.62										
NTZRM2001936	5.82	4.45	4.35	6.11	7.15	5.56	4.64	4.90	5.38										
NTZRM2001939	8.71	5.44	6.44	8.93	8.81	3.78	2.77	3.30	4.35										
NTZRM2001941	6.75	2.80	2.92	6.78	5.32	3.44	5.9	3.69	5.46										
NTZRM2001980	7.11	3.51	4.45	5.50	5.26	4.20	5.45	4.64	5.47										
NTZRM2001982	2.47	1.60	2.55	2.69	4.21	2.27	1.88	1.01	2.57										
NTZRM2001976	28.42	15.82	19.71	28.96	35.93	24.29	16.42	13.99	23.68										
NTZRM2001981	4.42	1.68	2.40	3.83	3.46	2.37	2.4	2.21	2.73										
NTZRM2001983	2.90	2.45	2.37	3.29	3.84	2.68	3.58	3.72	3.62										
NTZRM2001984	9.80	5.19	8.10	8.76	9.27	5.57	9.18	6.75	8.16										
NTZRM2001989	11.11	6.20	6.87	11.27	9.42	7.93	6.29	5.35	7.09										
NTZRM2001996	14.80	9.47	8.75	13.23	9.98	7.81	6.58	6.93	7.66										
NTZRM2001997	6.28	4.07	2.81	7.04	8.03	5.28	7.41	5.47	7.79										
NTZRM2001998	4.75	3.45	3.00	4.75	6.36	4.13	5.37	3.71	5.85										
NTZRM2001999	10.41	5.56	7.08	6.38	11.36	7.48	5.73	5.79	10.27										
NTZRM2002003	10.66	5.49	8.27	9.09	11.29	8.39	10.04	6.40	24.73										
NTZRM2002004	1.63	1.64	2.11	1.99	1.63	1.85	1.23	1.86	1.25										
NTZRM2002009	4.47	4.69	3.31	8.66	11.16	6.73	5.88	6.79	8.4	*	+								
NTZRM2002014	2.01	1.63	2.37	3.01	3.07	2.13	1.7	1.98	2.36										
NTZRM2002019	24.72	12.04	19.28	13.08	13.17	13.22	11.49	8.63	11.15										
NTZRM2002020	6.40	7.22	6.06	8.84	11.57	6.10	8.68	6.47	10.53										
NTZRM2002030	5.25	5.14	4.68	5.36	8.72	3.88	5.86	5.43	6.29										
NTZRM2002034	8.15	6.62	4.89	14.77	20.00	13.04	13.54	8.03	15.03	*	+								
NTZRM2002049	3.95	2.79	2.89	4.72	8.26	6.22	5.53	3.64	6.92	*	+								
NTZRM2002055	0.27	0.82	0.37	0.80	1.13	1.85	1.04	1.68	0.63										
NTZRM2002072	15.43	11.44	16.71	17.13	17.10	21.32	19.05	15.56	22.41										
NTZRM2002088	7.49	4.56	5.69	7.90	6.52	7.30	5.75	6.67	7.06										
NTZRM2002091	15.11	10.25	9.22	22.42	19.93	19.66	8.6	12.53	10.62	*	+								
NTZRM2002100	4.63	3.56	2.83	7.24	10.07	3.66	3.27	4.23	5.16										
NTZRM2002109	5.17	3.65	3.18	8.12	10.78	4.99	4.99	4.26	6.51										
NTZRM2002126	17.67	11.99	12.06	15.99	24.43	15.73	17.49	13.92	19.27										
NTZRM2002128	3.48	2.83	1.99	3.84	5.46	3.66	3.24	2.92	3.02										
NTZRM2002129	4.13	2.91	3.80	6.20	6.87	4.06	5.78	4.67	7.21										
NTZRM2002142	9.10	5.41	12.04	10.00	15.48	9.23	8.42	6.45	11.18										
NTZRM2002144	3.36	3.30	2.97	3.37	3.35	3.00	3.79	3.97	3.53										
NTZRM2002145	6.78	4.33	5.19	6.26	8.85	5.46	5.35	5.34	6.65										
NTZRM2002153	23.74	16.73	21.12	12.42	16.25	18.91	6.63	5.66	6.16										
NTZRM2002163	3.16	2.77	2.30	3.73	2.93	2.52	3.34	5.43	2.93										
NTZRM2002170	3.33	3.09	3.14	5.55	7.02	5.69	2.89	3.62	2.63	**	+								
NTZRM2002178	5.79	2.91	3.21	5.77	6.57	4.62	3.87	4.53	5.9										
NTZRM2002179	2.75	2.13	3.45	13.46	15.33	10.86	9.37	9.17	14.68	**	+								
NTZRM2002278	6.01	3.32	3.61	5.54	5.51	3.68	4.91	5.60	3.82										
NTZRM2002326	3.03	1.98	3.43	11.14	9.52	7.64	7.73	8.34	6.08	**	+								
NTZRM2002337	4.10	3.34	2.03	4.41	8.58	3.20	3.05	2.79	3.57										
NTZRM2002339	7.43	4.86	4.58	4.19	4.70	6.27	6.54	7.31	8.68										
NTZRM2002345	4.47	3.51	3.00	6.85	4.79	3.58	4.35	5.79	4.74										
NTZRM2002368	4.40	3.36	3.81	8.23	7.04	7.08	3.82	5.20	3.26	**	+								
NTZRM2002381	1.63	1.97	2.71	2.99	3.95	2.46	1.73	3.26	3.15										
NTZRM2002424	6.30	4.83	5.88	15.99	15.30	14.85	6.59	9.16	9.11	**	+								
NTZRM2002480	4.28	2.58	3.43	4.26	4.94	3.98	2.13	3.11	1.67										
NTZRM2002482	3.24	2.34	3.46	4.41	2.79	3.35	3.25	3.29	2.2										

1609

1610

NT2RM2002492	21.46	13.29	16.96	23.80	28.37	23.64	14.79	12.77	15.74	*	+		
NT2RM2002575	14.83	8.83	9.60	12.39	15.50	9.64	5.47	4.77	4.26		*	-	
NT2RM2002590	10.54	5.71	6.88	9.66	15.19	13.67	6.89	8.73	8.24				
NT2RM2002592	21.59	13.02	21.47	22.05	25.36	18.29	13.81	13.13	16.44				
NT2RM2002600	14.51	10.47	15.10	11.85	17.10	10.74	12.25	12.95	16.2				
NT2RM2002615	7.16	4.68	6.11	4.32	3.11	3.23	2.34	3.30	2.9	*	-	*	-
NT2RM2002622	7.42	4.82	9.06	37.13	40.07	28.33	10.87	12.05	11.06	**	+	*	+
NT2RM2002636	7.98	5.03	5.96	13.25	13.42	12.82	6.17	6.79	6.95	**	+		
NT2RM2002634	5.03	2.59	3.78	7.49	9.33	4.95	4.93	3.29	2.99				
NT2RM2002646	23.59	12.83	21.14	22.24	21.50	17.33	18.84	24.20	13.44				
NT2RM2002646	14.00	9.34	10.97	13.76	16.22	12.07	10.73	9.69	15.13				
NT2RM2002647	20.09	9.61	14.48	15.78	21.02	13.76	11.26	13.21	13.26				
NT2RM2002652	3.04	3.66	3.21	6.10	6.51	3.39	2.63	3.93	4.06				
NT2RM2002692	7.77	5.58	7.47	11.71	20.77	13.98	11.29	13.53	12.54	*	+	**	+
NT2RM2002721	24.72	15.21	18.70	28.40	46.33	25.49	20.79	20.54	32.86				
NT2RM2002748	79.54	53.04	79.10	79.94	73.77	78.90	26.04	31.33	37.42		*	-	
NT2RM2002764	5.43	3.03	2.52	10.76	7.77	4.58	3.86	3.77	3.77				
NT2RM2002772	11.93	7.88	8.81	11.61	12.84	7.73	4.61	5.99	7.99				
NT2RM2002811	9.63	5.90	5.86	8.67	8.08	6.76	5.99	5.21	6.14				
NT2RM2002818	6.94	3.95	3.88	7.36	7.54	5.33	2.65	4.06	4.31				
NT2RM2002879	2.57	1.77	2.32	2.29	3.75	1.84	3.18	4.11	4.37		*	+	
NT2RM2002979	11.80	7.84	8.67	10.47	13.00	9.87	8.38	6.63	6.92				
NT2RM2002981	4.73	2.96	3.25	4.20	5.55	4.27	4.3	5.20	4.19				
NT2RM2002995	3.40	2.64	2.64	3.84	3.50	4.10	2.62	3.34	2.85	*	+		
NT2RM2003031	3.92	1.02	1.63	4.33	4.68	2.72	3.7	2.74	3.72				
NT2RM2003042	21.41	10.74	8.21	17.59	19.62	15.87	7.89	8.90	9.64				
NT2RM2003044	3.74	2.06	1.81	3.99	6.41	3.64	2.33	3.97	3.12				
NT2RM2003099	4.60	2.18	1.89	2.49	4.89	3.16	3.07	3.31	2.92				
NT2RM2003096	3.67	1.54	1.20	3.30	4.47	3.32	3.18	3.65	3.25				
NT2RM2003116	5.36	5.13	6.83	5.86	7.80	6.25	3.24	6.72	6.31				
NT2RM2003222	2.53	2.08	1.54	2.39	2.31	1.74	0.73	3.10	1.35				
NT2RM2003224	15.53	10.87	13.94	24.44	25.63	15.64	6.09	8.22	11.35				
NT2RM2003250	14.48	5.65	5.15	9.14	10.21	4.29	3.99	3.24	3.21				
NT2RM2003250	2.29	2.33	1.33	2.70	2.97	1.92	4.64	2.60	3.37				
NT2RM2003262	12.60	10.45	8.76	10.06	13.00	11.50	9.36	7.15	7.82				
NT2RM4000023	1.99	1.44	1.54	4.90	4.52	3.88	4.13	2.29	4.66	**	+		
NT2RM4000024	2.91	2.48	1.20	3.30	4.50	2.17	2.67	1.90	2.29				
NT2RM4000027	8.33	4.07	5.06	2.82	3.04	1.62	1.79	2.08	2.61				
NT2RM4000030	5.84	5.94	5.16	8.87	6.03	4.15	5.42	5.51	5.41				
NT2RM4000033	1.51	1.27	1.03	2.93	3.16	1.42	1.59	1.08	1.27				
NT2RM4000034	2.39	1.22	1.22	3.53	2.94	1.45	2.28	1.04	1.5				
NT2RM4000046	2.68	1.77	1.53	3.42	3.11	1.75	3.04	1.82	3.01				
NT2RM4000052	4.15	1.71	1.72	3.48	3.49	2.40	3.28	1.71	3.37				
NT2RM4000054	26.80	19.29	17.31	21.55	22.04	23.11	25.09	20.51	27.5				
NT2RM4000061	2.10	1.10	0.99	1.68	1.71	1.22	2.51	1.77	1.98				
NT2RM4000074	9.55	7.34	6.67	13.37	15.17	7.83	6.11	5.94	7.27				
NT2RM4000085	2.96	0.88	2.51	4.65	4.96	3.33	2.05	2.94	4.07				
NT2RM4000086	5.73	3.89	4.54	5.27	5.33	3.12	1.63	3.66	5.45				
NT2RM4000100	5.36	2.82	2.66	5.25	5.01	3.76	5.82	4.53	4.32				
NT2RM4000101	3.85	2.50	2.70	3.14	2.31	2.97	4.25	3.04	4.67				
NT2RM4000102	36.64	21.10	21.71	40.33	40.26	40.80	33.11	25.16	36.34				
NT2RM4000104	1.41	0.89	0.77	2.16	1.98	1.38	2.39	2.42	1.99		**	+	
NT2RM4000115	1.25	1.28	1.23	1.59	2.32	1.33	1.87	1.24	1.33				
NT2RM4000129	2.55	2.06	1.92	3.48	3.51	3.33	3.04	3.19	2.54	**	+		
NT2RM4000139	2.48	1.32	1.75	2.52	1.95	2.96	1.63	2.77	2.58				
NT2RM4000149	1.92	1.98	1.88	3.18	3.67	2.07	1.43	1.95	2.57				

67/180

【0708】

【表237】

1611

1612

NT2RM4000154	8.41	4.25	5.83	5.71	7.89	3.63	6.31	2.89	10.88										
NT2RM4000156	4.06	2.82	3.12	3.91	5.14	3.25	4.15	3.34	7.54										
NT2RM4000167	2.76	1.86	2.44	3.27	3.78	2.46	1.7	2.54	2.08										
NT2RM4000169	19.79	11.82	12.59	15.78	28.83	16.15	10.62	8.62	22.74										
NT2RM4000191	5.46	2.93	3.98	7.00	12.87	3.95	5.75	4.22	5										
NT2RM4000197	6.21	3.61	5.57	1.78	3.32	3.20	2.07	2.72	3.62										
NT2RM4000198	6.32	5.24	5.02	9.16	10.86	8.33	6.38	7.74	6.83	**	+								
NT2RM4000199	3.97	1.83	1.79	3.99	4.05	3.81	2.77	3.05	3.55										
NT2RM4000208	3.35	2.42	1.54	4.45	2.14	1.95	1.94	2.20	2.16										
NT2RM4000202	3.63	1.09	1.43	2.56	2.87	2.44	2.2	2.07	1.78										
NT2RM4000218	4.14	2.52	2.72	3.86	8.22	3.80	3.01	2.97	3.68										
NT2RM4000215	5.18	3.07	5.47	7.27	8.43	5.15	4.83	4.89	4.29										
NT2RM4000220	2.94	2.54	2.79	4.64	4.57	3.49	4.1	4.60	5.77	*	+	*	+						
NT2RM4000229	5.01	3.09	3.00	5.45	4.41	4.69	4.07	4.71	4.56										
NT2RM4000231	4.55	4.22	5.24	5.48	9.85	6.48	5.25	5.36	6.29										
NT2RM4000233	15.69	9.94	12.92	10.36	8.30	6.63	11.95	12.79	13.03										
NT2RM4000244	3.55	2.12	1.68	2.06	1.74	1.35	2.28	2.40	1.4										
NT2RM4000251	3.33	1.28	1.28	2.48	6.47	3.24	2.39	3.65	3.7										
NT2RM4000255	2.86	2.95	2.55	3.65	4.00	4.45	3.46	3.56	4.05	**	+	**	+						
NT2RM4000265	4.79	2.78	4.25	9.35	12.26	8.62	3.89	4.46	7.5	**	+								
NT2RM4000283	70.67	47.66	58.69	22.90	27.64	23.33	20.04	20.53	29.33	**	-	**	-						
NT2RM4000284	3.79	2.43	3.13	4.73	5.37	4.18	3.75	4.06	5.01	*	+								
NT2RM4000290	3.63	2.15	2.31	4.25	6.01	4.45	4.22	4.40	5.11	*	+	*	+						
NT2RM4000295	2.18	1.74	1.84	1.64	1.85	1.54	2.16	2.51	2.05										
NT2RM4000306	9.76	5.69	5.53	3.29	5.79	3.80	4.99	4.91	4.19										
NT2RM4000307	1.99	1.95	1.34	6.27	6.75	5.25	9.66	12.35	13.1	**	+	**	+						
NT2RM4000309	4.39	2.45	3.20	3.45	3.57	3.25	2.21	2.77	3.12										
NT2RM4000313	4.53	2.93	3.37	6.76	7.38	6.57	4.37	4.56	4.95	**	+								
NT2RM4000318	3.24	1.42	3.10	6.35	5.08	6.14	3.2	4.49	3.95	**	+								
NT2RM4000324	3.33	2.91	2.72	5.10	4.10	4.09	3.41	4.13	3.13	*	+								
NT2RM4000326	2.66	2.08	2.02	2.52	2.48	2.90	1.91	4.16	2.37										
NT2RM4000327	5.98	3.83	5.87	11.13	9.36	9.04	5.82	4.08	6.84	**	+								
NT2RM4000344	18.32	6.89	6.35	13.95	16.21	14.72	10.48	11.38	12.84										
NT2RM4000349	6.58	3.84	3.66	6.40	5.99	6.38	4.94	4.61	4.8										
NT2RM4000354	5.00	2.70	3.97	3.28	2.86	2.19	2.4	2.57	3.45										
NT2RM4000356	4.16	1.61	1.73	2.39	4.18	5.03	2.81	3.86	2.62										
NT2RM4000366	51.05	23.81	40.37	61.56	72.80	50.45	36.85	39.74	37.86										
NT2RM4000368	4.89	2.95	4.56	12.45	6.89	8.75	3.93	5.00	5.04	*	+								
NT2RM4000372	3.91	2.54	3.44	5.84	6.63	5.55	3.15	4.00	4.07	**	+								
NT2RM4000386	2.58	1.67	2.32	2.56	2.07	2.16	1.54	2.11	1.77										
NT2RM4000395	7.43	3.02	3.38	5.38	8.33	4.62	3.75	3.62	2.41										
NT2RM4000414	8.01	4.62	4.45	4.72	6.23	4.47	6.27	6.74	7.44										
NT2RM4000417	3.81	2.15	2.35	2.45	4.44	3.37	1.96	2.94	5.43										
NT2RM4000431	4.32	3.14	3.21	5.52	5.35	4.92	3.28	3.80	2.75	*	+								
NT2RM4000435	5.83	3.82	4.77	12.15	12.72	12.65	6.8	7.68	8.72	**	+	*	+						
NT2RM4000433	3.24	1.87	2.39	3.27	3.60	3.54	3.12	3.76	4.32			*	+						
NT2RM4000436	5.20	2.98	5.09	4.80	5.70	3.50	3.27	3.38	2.58										
NT2RM4000444	2.77	3.48	2.67	4.83	3.05	2.47	3.64	2.65	2.6										
NT2RM4000457	15.74	6.60	7.42	19.58	21.46	13.99	7.15	6.46	8.49										
NT2RM4000471	2.61	2.36	2.45	4.93	5.40	4.25	2.75	2.88	3.1	**	+	*	+						
NT2RM4000472	18.08	9.02	12.03	39.03	47.24	22.44	10.92	7.97	22.46	*	+								
NT2RM4000486	3.65	3.27	2.92	5.46	7.04	5.93	3.74	4.42	3.79	**	+								
NT2RM4000490	4.88	4.19	2.87	2.63	4.80	4.02	3.72	6.17	4.86										
NT2RM4000495	4.08	3.13	4.22	2.96	3.44	3.46	3.56	4.42	3.51										
NT2RM4000505	13.63	8.59	11.63	16.76	17.74	14.63	12.93	15.48	13.33	*	+								
NT2RM4000511	58.96	34.63	49.12	52.44	54.53	48.10	20.11	21.96	24.23			*	-						

68/180

[0709]

[表238]

1614

N7ZRM4000514	5.53	2.38	2.75	8.23	11.94	5.81	3.95	5.11	3.73										
N7ZRM4000515	16.72	6.51	7.89	17.68	18.19	15.60	8.65	8.97	10.58										
N7ZRM4000517	52.07	29.36	32.53	47.60	48.78	40.92	19.63	19.22	17.95										
N7ZRM4000520	2.37	1.45	1.44	1.17	1.70	1.58	0.83	2.01	2.17										
N7ZRM4000531	1.99	2.27	1.67	2.66	3.68	3.90	3.09	4.12	3.64	*									
N7ZRM4000532	1.32	0.65	0.82	1.96	2.81	1.58	1.14	2.83	2.21	*	+								
N7ZRM4000533	3.05	2.29	3.20	1.70	2.71	1.77	1.32	2.54	1.44										
N7ZRM4000534	1.94	0.89	1.21	1.63	2.79	1.54	1.47	2.29	1.5										
N7ZRM4000563	8.72	3.55	3.49	6.44	4.79	3.51	5.01	4.09	5.24										
N7ZRM4000566	4.57	2.22	2.28	4.38	4.92	2.84	2.28	2.65	3.1										
N7ZRM4000568	3.97	2.58	1.85	3.65	4.45	3.11	2.68	3.32	5.31										
N7ZRM4000585	4.60	2.16	1.71	2.71	3.64	3.29	2.11	2.49	3.12										
N7ZRM4000587	2.44	1.07	2.17	2.90	3.56	2.74	2.55	3.03	3.48										
N7ZRM4000590	2.10	1.53	1.91	1.79	3.35	2.47	1.66	3.17	1.73										
N7ZRM4000593	7.87	4.39	5.71	12.59	12.23	10.68	4.27	7.40	6.4	**	+								
N7ZRM4000595	2.17	1.55	2.08	3.28	3.82	2.26	2.16	3.13	4.82										
N7ZRM4000603	12.55	6.66	4.52	9.64	5.73	7.27	4.44	2.74	3.84										
N7ZRM4000611	4.28	4.40	1.85	3.49	3.25	3.20	4.67	2.74	3.19										
N7ZRM4000616	3.34	2.92	3.77	4.32	4.33	3.69	3.56	2.86	2.97										
N7ZRM4000621	16.48	12.72	9.94	21.48	22.15	19.49	9.54	8.56	8.57	*	+								
N7ZRM4000648	2.01	1.43	1.11	2.32	1.95	1.99	1.76	2.62	1.65										
N7ZRM4000649	5.47	3.71	4.22	6.21	6.35	6.84	6.07	5.86	5.42	*	+								
N7ZRM4000658	8.60	4.07	5.16	8.70	7.92	4.74	5.84	5.98	5.36										
N7ZRM4000661	10.99	4.92	5.69	11.11	10.38	8.21	15.64	14.68	17.57										
N7ZRM4000673	9.96	5.23	4.31	6.63	5.66	5.28	8.2	5.95	5.83										
N7ZRM4000674	5.01	2.88	2.93	4.58	4.03	4.02	5.28	3.25	4.19										
N7ZRM4000689	6.44	3.20	3.50	4.50	6.19	4.47	3.52	4.05	3.79										
N7ZRM4000698	35.87	22.93	21.16	15.46	17.90	22.28	17.5	16.82	14.8										
N7ZRM4000700	3.46	2.08	2.83	3.85	2.02	2.52	2.49	2.37	1.92										
N7ZRM4000701	9.78	5.90	5.74	10.46	14.71	8.86	7.95	6.35	8.32										
N7ZRM4000712	2.69	1.64	2.42	4.68	4.33	3.64	2.57	3.33	2.41	*	+								
N7ZRM4000717	12.02	5.07	6.36	11.87	8.62	8.11	7.27	6.28	7.15										
N7ZRM4000733	8.98	3.57	6.27	6.72	6.26	7.78	7.76	4.90	6										
N7ZRM4000734	9.72	3.11	3.90	7.75	4.13	5.58	5.8	4.00	5.07										
N7ZRM4000741	4.49	2.29	3.36	3.14	3.42	3.32	3.44	4.03	2.18										
N7ZRM4000744	3.69	2.68	2.61	2.80	6.32	4.46	2.85	3.92	3										
N7ZRM4000749	11.40	7.45	11.83	11.62	13.08	12.36	13.08	12.48	13.4										
N7ZRM4000751	6.54	4.81	4.52	15.28	14.53	10.59	6.43	6.81	9.13	**	+								
N7ZRM4000752	4.53	2.37	3.48	4.41	5.68	4.78	3.23	4.75	8.68										
N7ZRM4000760	4.53	2.84	2.99	5.14	6.37	2.91	5.41	3.73	5.34										
N7ZRM4000761	996.52	787.70	799.46	925.45	928.68	688.98	521.4	1076.26	1043										
N7ZRM4000764	27.63	19.80	15.48	20.84	20.29	16.92	30.21	26.08	33.56										
N7ZRM4000768	14.67	8.26	9.77	8.91	9.00	6.52	3.2	6.21	5.06										
N7ZRM4000778	4.92	2.41	4.01	2.84	3.65	2.97	1.85	2.67	2.07										
N7ZRM4000779	8.60	6.98	9.29	9.01	13.32	14.40	9.71	7.63	14.65										
N7ZRM4000787	4.24	2.50	3.69	7.64	7.50	6.95	5.13	4.57	3.51	**	+								
N7ZRM4000790	3.29	2.32	3.49	4.70	4.95	5.71	2.8	3.89	2.61	*	+								
N7ZRM4000795	17.99	8.62	8.95	7.60	7.29	5.12	10.59	11.21	13.05										
N7ZRM4000796	9.52	5.97	4.89	6.98	7.91	6.65	7.34	5.94	6.5										
N7ZRM4000798	4.86	3.32	1.92	4.08	3.21	6.07	3.4	3.16	3.56										
N7ZRM4000800	25.53	16.14	15.27	24.04	32.78	23.66	18.49	15.32	20.57										
N7ZRM4000813	2.68	4.14	3.79	3.70	5.76	3.65	5.28	6.71	6.8										
N7ZRM4000820	6.65	4.53	5.35	8.29	7.69	8.43	5.66	5.99	4.55	*	+								
N7ZRM4000827	7.32	3.89	5.09	8.78	8.63	9.18	6.29	5.93	5.97	*	+								
N7ZRM4000830	6.10	3.43	4.84	5.47	6.76	7.08	5.66	6.54	4.35										
N7ZRM4000833	7.52	4.61	4.22	4.98	5.08	4.81	5.7	5.33	4.23										

1616

N1ZRM4008841	5.06	3.39	2.43	4.05	5.93	4.27	4.63	4.44	4.32				
N1ZRM4008846	9.09	5.94	7.28	12.84	12.70	15.11	9.96	10.78	11.36	**	+	+	+
N1ZRM4008848	7.88	5.40	5.25	6.98	11.06	5.33	7.53	6.82	8.25				
N1ZRM4008853	6.75	4.64	5.34	13.69	17.70	14.08	11.97	13.34	10.93	**	+	**	+
N1ZRM4008855	4.73	2.86	4.28	6.84	5.05	6.75	4.95	4.71	3.45		+		
N1ZRM4008859	19.33	7.63	8.66	12.33	11.71	13.85	10.92	13.05	13.48				
N1ZRM4008868	3.39	2.48	3.24	2.56	3.27	2.72	2.54	2.52	2.34				
N1ZRM4008878	7.43	4.59	4.58	4.56	7.18	4.83	5.21	5.55	10.16				
N1ZRM4008879	5.36	4.71	2.54	2.94	5.60	3.69	4.73	3.05	8.38				
N1ZRM4008882	13.28	7.67	8.34	13.87	16.02	12.84	11.37	9.53	8.64				
N1ZRM4008887	7.73	5.89	6.66	6.98	5.77	6.42	10.56	10.15	7.39				
N1ZRM4008895	5.72	3.47	4.08	7.64	7.37	6.94	4.46	6.14	5.93		+		
N1ZRM4008897	7.53	4.28	4.64	7.90	11.04	6.20	7.51	8.32	7.28				
N1ZRM40089901	2.04	1.85	1.79	2.60	2.63	3.31	2.13	2.92	1.47	*			
N1ZRM4008950	0.56	0.78	1.17	2.14	1.27	1.24	1.41	2.19	1.17				
N1ZRM4008965	9.86	4.20	4.55	3.72	5.50	4.12	5.03	3.46	4.87				
N1ZRM4008971	5.30	5.00	2.48	7.54	6.04	2.89	3.53	4.64	7.17				
N1ZRM4008979	4.99	2.53	1.69	2.02	3.14	2.85	2.38	2.83	3.27				
N1ZRM4008987	2.44	1.53	2.68	3.20	3.68	2.75	2.62	4.83	3.64				
N1ZRM4008989	4.94	3.38	3.37	4.04	2.94	2.51	3.27	4.13	3.58				
N1ZRM4008991	0.93	1.02	1.31	2.15	2.31	2.55	2.33	4.87	2.11	**	+		
N1ZRM4008992	11.24	7.63	10.16	7.25	5.43	5.90	4.54	4.54	4.18	*	-	**	
N1ZRM4008996	4.06	2.34	3.75	9.54	9.91	8.12	3.46	4.48	3.87	**	+		
N1ZRM4008997	9.49	3.35	2.92	6.90	7.64	7.96	5.25	6.12	5.29				
N1ZRM4001001	22.10	15.26	10.21	12.02	9.69	11.49	22.6	17.92	9.97				
N1ZRM4001002	5.24	3.19	3.25	8.21	8.99	8.70	5.14	6.05	8.69	**			
N1ZRM4001016	4.56	3.14	3.04	3.93	5.46	2.92	3.16	3.93	3.9				
N1ZRM4001025	115.98	53.32	70.45	58.33	60.27	42.54	40.15	40.87	41.74				
N1ZRM4001027	0.14	0.43	0.68	0.22	0.51	0.94	0.68	1.67	1.36				
N1ZRM4001032	1.80	1.46	0.81	3.10	2.87	2.32	1.9	2.71	1.77	**	+		
N1ZRM4001047	1.37	0.85	0.95	2.05	2.61	2.62	1.72	2.01	1.51	**	+	+	+
N1ZRM4001049	10.71	3.63	3.82	6.40	6.54	4.49	5.52	5.09	5.26				
N1ZRM4001051	6.70	3.93	4.20	7.11	12.15	5.54	5.61	4.11	11.9				
N1ZRM4001053	8.14	4.27	4.08	6.07	7.39	5.45	8.57	7.89	6.02				
N1ZRM4001053	27.19	14.20	21.35	17.33	19.31	15.07	12.02	9.63	10.5				
N1ZRM4001054	3.61	1.72	2.96	2.73	3.57	4.09	2.66	3.55	3.62				
N1ZRM4001059	7.61	4.52	5.00	8.40	9.15	6.24	6.45	6.67	1.85				
N1ZRM4001071	4.06	2.69	2.57	4.40	6.02	4.14	3.25	5.00	2.66				
N1ZRM4001084	4.94	2.76	3.04	3.73	6.39	5.46	4.17	4.56	4.31				
N1ZRM4001092	7.29	2.48	2.72	5.06	4.22	4.55	3.22	2.32	2.04				
N1ZRM4001100	12.18	6.64	7.67	10.87	11.09	10.86	6.95	8.94	8.4				
N1ZRM4001116	1.86	1.58	1.69	2.27	2.62	2.03	2.58	1.98	1.6	**	+		
N1ZRM4001119	4.12	2.84	2.77	3.79	5.02	3.34	2.73	3.61	4.07				
N1ZRM4001148	16.77	10.70	11.39	11.80	11.74	11.76	7	6.89	6.74			*	-
N1ZRM4001148	13.85	6.50	6.41	8.02	8.87	5.20	9.72	12.16	8.38				
N1ZRM4001151	3.04	2.82	2.68	3.38	3.91	4.17	3.34	5.07	4.04	*	+		
N1ZRM4001155	3.85	1.95	2.51	2.48	2.96	3.06	2.88	3.51	1.43				
N1ZRM4001157	4.58	2.01	1.48	3.42	3.84	2.43	3.68	3.71	2.97				
N1ZRM4001160	6.16	2.57	2.15	5.06	4.60	3.14	2.68	2.65	4.39				
N1ZRM4001163	28.46	18.93	15.20	35.93	37.53	27.65	20.27	18.39	15.85				
N1ZRM4001187	5.13	3.42	2.71	6.56	6.27	4.41	3.87	4.84	5.54				
N1ZRM4001191	4.08	1.58	2.81	4.80	3.69	3.67	1.67	2.71	2.13				
N1ZRM4001200	5.87	3.25	4.14	11.90	10.51	10.62	3.7	6.92	6.13	**	+		
N1ZRM4001203	5.49	3.54	4.23	3.75	6.16	5.89	3.38	6.37	4.68				
N1ZRM4001204	1.21	0.66	1.10	1.28	1.38	1.01	0.49	2.21	0.81				
N1ZRM4001217	2.79	2.05	1.40	2.03	2.19	2.27	3.13	2.64	2.28				

1617

1618

NT2RM4001245	8.44	3.64	3.02	4.82	4.98	4.20	6.1	4.52	4.33				
NT2RM4001247	3.08	1.77	1.70	4.91	4.46	4.61	4.28	3.78	3.75	**	+	*	+
NT2RM4001256	2.68	1.97	1.44	2.55	3.37	2.58	2.83	3.46	2.47				
NT2RM4001258	3.01	1.08	1.34	2.58	2.80	3.08	2.91	3.65	2.07				
NT2RM4001267	3.85	1.81	3.09	2.74	2.85	2.20	1.95	2.54	1.48				
NT2RM4001273	4.22	3.00	2.18	5.27	4.13	4.07	4.07	4.58	3.5				
NT2RM4001281	4.83	2.17	2.72	3.21	3.18	3.70	4.92	3.04	3.76				
NT2RM4001286	200.90	135.14	135.42	284.75	209.56	246.97	164.2	134.29	151.3	*	+		
NT2RM4001290	9.86	4.80	5.69	5.57	5.18	5.39	8.08	8.05	9.32				
NT2RM4001309	4.86	3.06	2.25	4.98	6.28	4.18	3.55	4.91	3.92				
NT2RM4001313	5.02	3.13	3.38	10.23	11.21	8.30	5.64	5.09	6.07	**	+		
NT2RM4001316	3.10	1.87	1.63	4.90	3.32	2.72	2.34	3.07	2.48				
NT2RM4001320	3.57	1.99	1.80	4.35	3.25	2.99	2.67	3.38	1.95				
NT2RM4001321	2.36	1.76	2.19	4.88	3.23	3.63	2.96	3.26	2.18	*	+		
NT2RM4001325	4.26	2.86	2.43	3.61	4.06	3.37	3.66	2.87	3.97				
NT2RM4001333	9.63	4.30	7.26	19.73	18.36	12.94	10.99	11.48	14.86	*	+		
NT2RM4001340	15.08	7.81	6.58	8.67	7.96	8.93	6.1	7.09	9.47				
NT2RM4001344	5.69	1.98	2.69	4.58	3.47	5.21	3.57	4.25	3.42				
NT2RM4001347	2.27	2.16	1.78	2.66	5.15	3.34	3.4	3.43	2.43			*	+
NT2RM4001357	6.72	4.15	5.35	6.32	6.10	5.55	4.34	5.12	6.64				
NT2RM4001360	5.77	3.29	3.38	4.26	4.44	4.12	4.69	3.72	3.64				
NT2RM4001371	4.54	2.79	3.83	7.15	6.45	5.83	3.62	4.03	2.04	*	+		
NT2RM4001377	10.12	5.47	3.83	5.72	6.90	5.90	6.53	6.36	7.54				
NT2RM4001382	27.64	18.16	15.30	26.18	25.29	24.42	17.41	14.13	18.42				
NT2RM4001384	2.18	1.75	1.21	2.08	4.07	2.57	1.75	1.84	2.63				
NT2RM4001400	1.97	1.68	1.05	5.11	4.43	3.04	4.16	3.64	2.67	*	+	*	+
NT2RM4001409	2.47	2.29	2.32	4.11	6.40	4.45	3.11	3.39	3.96	*	+	*	+
NT2RM4001410	3.95	1.97	3.57	4.82	7.04	5.31	4.02	3.43	5.37	*	+		
NT2RM4001411	0.83	0.77	0.89	2.84	2.80	2.65	2.26	2.50	1.14	**	+		
NT2RM4001413	3.72	2.65	2.59	3.20	3.12	4.78	3.05	4.81	2.12				
NT2RM4001414	4.96	2.76	1.91	3.88	3.24	3.95	8.58	4.11	4.61				
NT2RM4001436	10.71	5.74	4.93	8.68	8.18	5.45	5.99	5.69	6.31				
NT2RM4001437	3.31	2.10	1.69	4.84	3.86	4.25	2.81	3.12	5.09	*	+		
NT2RM4001444	17.08	11.93	9.02	14.34	23.31	17.28	9.91	10.57	15.43				
NT2RM4001454	1.52	1.27	0.92	2.66	2.43	3.18	2.53	2.75	3.91	**	+	*	+
NT2RM4001455	1.97	1.35	0.94	1.41	2.43	2.26	1.92	2.49	2.53				
NT2RM4001483	8.15	6.48	6.83	17.59	20.73	16.59	7.89	9.09	9.1	**	+		
NT2RM4001489	2.71	2.11	2.58	4.94	4.32	3.30	3.82	3.97	2.42	*	+		
NT2RM4001495	18.14	8.14	7.60	6.61	8.27	8.97	13.02	9.27	7.52				
NT2RM4001499	12.77	8.16	6.92	3.39	3.00	2.48	3.08	3.42	2.67	*	-	*	-
NT2RM4001515	3.27	1.91	1.68	2.35	4.06	1.83	1.52	2.44	1.37				
NT2RM4001519	5.12	2.84	4.04	2.41	3.33	2.32	2.38	4.57	1.4				
NT2RM4001522	6.04	4.16	3.86	10.17	8.78	6.98	5.57	5.11	4.64	*	+		
NT2RM4001523	2.87	2.23	1.80	2.40	4.75	2.55	2.53	3.39	1.48				
NT2RM4001550	9.31	4.21	5.82	7.65	10.18	9.65	4.79	5.78	4.65				
NT2RM4001553	13.10	6.91	9.72	15.17	15.42	12.48	9.84	10.30	8.03				
NT2RM4001554	6.26	1.91	2.23	3.10	3.46	2.19	2.03	3.40	3.47				
NT2RM4001557	1.82	1.50	1.72	2.44	4.16	3.37	2.15	2.77	2.22	*	+	*	+
NT2RM4001565	4.45	2.55	3.09	4.16	3.19	4.16	3.34	4.45	3.44				
NT2RM4001566	8.15	6.36	5.54	21.07	22.32	19.38	14.82	13.59	12.47	**	+	**	+
NT2RM4001569	1.07	2.72	1.12	1.58	1.44	1.53	1.39	2.06	0.92				
NT2RM4001579	2.12	1.63	1.82	2.74	2.69	2.72	4.53	3.33	2.15	**	+		
NT2RM4001582	2.62	2.33	2.55	3.71	4.48	4.20	3.06	3.87	3.26	**	+	*	+
NT2RM4001589	8.35	5.09	6.66	12.13	12.37	10.11	11.51	12.65	14.42	*	+	**	+
NT2RM4001522	3.41	2.19	1.04	2.79	0.97	1.51	1.07	1.30	2.99				
NT2RM4001594	6.13	3.39	4.24	4.36	6.50	3.95	3.79	5.84					

71/180

【0712】

【表241】

1619

1620

NT2RM4001597	9.12	5.34	5.09	9.11	10.92	8.47	7.88	8.77	8.89										
NT2RM4001605	2.56	1.50	0.61	1.85	2.19	2.01	1.99	3.21	1.69										
NT2RM4001609	89.25	51.45	54.24	71.13	77.23	52.58	39.95	36.11	41.37										
NT2RM4001610	12.00	8.23	7.07	12.20	9.44	8.76	11.99	11.53	14.72										
NT2RM4001611	2.42	1.85	2.60	3.39	3.10	2.31	2.05	3.53	1.91										
NT2RM4001618	9.99	6.27	7.80	11.85	10.16	11.99	7.45	6.31	7.61										
NT2RM4001622	26.67	8.64	17.82	10.07	12.08	11.47	11.1	11.92	5.45										
NT2RM4001624	6.68	3.27	2.64	4.78	7.08	4.67	4.35	5.33	5.32										
NT2RM4001625	6.46	4.15	3.63	6.09	6.98	6.57	5.81	6.49	4.68										
NT2RM4001629	3.08	1.43	1.44	3.13	3.87	3.98	3.34	3.46	2.65										
NT2RM4001633	29.86	24.78	26.14	43.08	46.42	34.45	16.71	16.75	13.76	*	+	**	-						
NT2RM4001642	2.85	2.24	1.81	3.57	2.70	1.88	1.79	3.45	2.28										
NT2RM4001647	17.28	7.78	9.99	11.15	12.30	10.77	8	8.01	6.38										
NT2RM4001659	0.99	1.51	1.38	2.58	3.80	3.02	1.93	2.32	1.3	**	+								
NT2RM4001662	7.87	3.75	2.87	5.79	6.00	4.16	5.7	4.40	5.34										
NT2RM4001666	5.31	2.73	1.99	5.11	5.72	2.91	2.77	3.37	5										
NT2RM4001670	11.64	5.63	4.93	10.66	7.77	4.83	7.89	5.98	5.85										
NT2RM4001682	7.63	4.69	7.88	11.61	13.13	10.67	7.98	7.62	9.49	*	+								
NT2RM4001710	3.51	1.93	3.14	2.89	2.81	2.52	2.94	3.14	3.23										
NT2RM4001712	4.09	1.48	2.36	6.28	6.47	3.67	3.14	2.79	2.86										
NT2RM4001714	9.74	6.27	6.28	8.33	6.94	5.10	4.33	4.54	3.78										
NT2RM4001715	9.70	6.79	8.58	10.69	5.46	8.50	6.49	7.88	6.36										
NT2RM4001727	9.24	3.95	4.64	8.67	8.28	6.42	5.55	4.51	4.54										
NT2RM4001731	13.05	6.04	4.43	9.34	11.19	3.94	6.46	7.94	7.44										
NT2RM4001735	10.60	7.33	6.23	6.67	8.99	10.11	4.77	6.71	9.86										
NT2RM4001739	4.78	4.21	5.14	4.57	4.78	3.04	2.46	4.65	3.94										
NT2RM4001741	9.97	6.74	4.99	10.67	11.48	8.89	9.93	7.28	7.04										
NT2RM4001746	4.40	2.92	3.08	6.46	6.23	6.82	4.23	5.87	3.98	**	+								
NT2RM4001754	5.88	4.22	4.77	3.77	2.85	3.40	2.26	3.95	2.51	*	-	*	-						
NT2RM4001757	3.98	2.34	2.64	6.30	5.38	5.11	4.27	5.17	3.56	*	+								
NT2RM4001758	4.03	1.40	1.41	2.95	3.14	0.90	2.11	1.49	2.63										
NT2RM4001768	9.33	3.18	2.78	8.73	9.23	6.03	4.74	5.46	7.46										
NT2RM4001775	1.60	0.85	0.48	1.68	1.19	1.13	0.51	1.89	2.16										
NT2RM4001776	1.24	0.67	0.70	2.08	1.65	1.01	0.84	1.25	1.26										
NT2RM4001783	3.30	1.81	1.77	3.52	4.08	2.55	1.62	3.51	1.6										
NT2RM4001793	5.58	4.64	4.50	8.16	8.15	6.01	4.19	4.76	4.23	*	+								
NT2RM4001810	3.48	2.21	2.29	3.20	3.69	2.65	2.04	3.39	2.03										
NT2RM4001813	3.11	0.62	1.16	2.31	2.18	1.56	2	3.91	2.71										
NT2RM4001818	3.22	2.40	2.49	5.46	4.70	3.11	4.89	3.44	5.14		*	+							
NT2RM4001819	11.19	5.78	6.63	9.35	9.42	7.47	10.81	7.51	7.34										
NT2RM4001823	3.13	1.86	1.29	2.61	3.40	2.11	3.37	1.94	1.66										
NT2RM4001828	8.26	6.14	6.03	15.07	18.35	11.62	9.17	6.53	11.85	*	+								
NT2RM4001835	3.34	2.52	2.50	5.07	6.41	5.16	6.93	7.44	8.93	**	+	**	+						
NT2RM4001836	3.42	2.60	1.50	3.55	5.57	1.89	3.02	2.62	2.83										
NT2RM4001841	7.03	4.07	5.20	3.69	3.84	5.28	4.46	6.00	6.72										
NT2RM4001842	2.54	1.03	0.84	4.40	5.14	3.61	2.15	3.10	2.2	*	+								
NT2RM4001843	7.33	3.08	3.29	4.61	4.36	4.19	6.63	4.29	4.74										
NT2RM4001856	7.28	3.36	2.92	6.92	6.61	6.34	6.76	4.92	39.96										
NT2RM4001858	4.41	2.01	2.89	4.19	5.25	3.77	3.99	3.14	3.55										
NT2RM4001861	15.16	9.14	7.90	8.10	8.14	2.12	7.69	7.66	6.31										
NT2RM4001863	5.18	5.03	4.89	5.35	5.57	4.84	4.1	2.95	4.25		*	-							
NT2RM4001865	4.40	1.50	1.71	4.54	5.77	6.01	3.87	4.69	4.27	*	+								
NT2RM4001869	6.80	4.12	4.66	5.90	4.78	4.71	3.79	3.46	3.8										
NT2RM4001873	9.91	7.88	7.75	6.45	7.32	6.28	5.39	4.87	5.18		**	-							
NT2RM4001876	20.13	9.94	9.70	9.48	8.26	10.48	13.84	12.07	14.41										
NT2RM4001880	6.36	4.04	3.70	6.23	5.32	5.66	5.53	4.83	6.28										

72/180

【0713】

【表242】

1621

1622

NT2RM4001885	12.23	5.39	5.31	15.89	14.89	14.46	9.96	8.97	11.11	*	+		
NT2RM4001889	17.90	10.90	9.56	25.74	24.82	26.44	14.72	12.91	12.79	**	+		
NT2RM4001894	3.99	3.32	3.07	4.15	4.34	4.16	5.09	3.83	3.49				
NT2RM4001897	4.68	3.36	3.66	5.57	7.84	6.03	9.17	7.60	6.62	*	+	**	+
NT2RM4001899	4.37	2.59	2.66	5.10	4.85	5.00	3.8	4.79	3.2	*	+		
NT2RM4001905	14.13	19.47	18.60	6.62	5.76	7.88	4.18	4.49	4.16	**	-	**	-
NT2RM4001921	4.57	2.06	2.67	5.98	6.27	5.24	3.2	3.09	2.6	*	+		
NT2RM4001930	7.89	5.36	5.01	6.12	7.65	5.79	3.76	3.60	3.88				
NT2RM4001938	3.35	3.03	2.31	4.03	4.25	3.01	4.12	3.88	3.78		*	+	
NT2RM4001940	8.88	7.21	7.25	7.65	9.61	6.94	5.41	5.68	5.3		*	-	
NT2RM4001943	48.53	24.69	36.33	81.10	98.59	62.39	57.37	65.30	79.98	*	+	*	+
NT2RM4001953	4.86	4.02	3.80	11.16	10.73	8.47	5.44	7.13	6.71	**	+	*	+
NT2RM4001965	3.95	3.09	2.78	3.89	4.20	5.02	3.08	4.34	1.87				
NT2RM4001966	4.92	2.59	2.69	5.18	4.42	3.96	3.32	4.68	3.49				
NT2RM4001969	4.52	3.56	2.88	4.01	4.54	3.26	3.65	2.05	3.76				
NT2RM4001974	3.18	2.93	2.68	3.45	3.46	4.29	4	3.93	2.9				
NT2RM4001979	7.10	5.28	4.65	8.51	9.51	9.19	5.57	5.12	5.65	*	+		
NT2RM4001988	8.43	6.53	5.48	9.14	11.80	9.30	5.72	6.09	7.18				
NT2RM4001984	0.37	0.36	2.68	1.04	2.24	1.27	3.83	2.41	1.54				
NT2RM4001987	5.43	3.22	4.46	5.44	5.41	4.74	6.11	4.65	5.13				
NT2RM4002013	4.01	2.99	3.04	5.45	6.17	4.31	4.16	6.39	4.96	*	+		
NT2RM4002018	1.35	1.30	1.91	4.17	2.80	1.86	2.66	3.82	2.52		*	+	
NT2RM4002033	5.95	4.44	3.94	8.70	9.58	8.70	6.99	4.97	5.08	**	+		
NT2RM4002034	10.16	6.70	5.00	9.69	8.87	7.70	7.22	5.62	6.43				
NT2RM4002044	17.29	9.91	9.34	16.54	14.23	14.16	9.93	9.20	9.33				
NT2RM4002047	4.89	3.52	4.39	7.70	9.18	8.38	5.94	5.42	6.2	**	+	*	+
NT2RM4002054	5.22	3.24	3.62	4.72	4.27	3.95	3.64	4.57	3.02				
NT2RM4002055	4.93	3.27	3.62	3.58	4.71	3.15	4.03	4.74	4.4				
NT2RM4002059	10.05	6.75	9.67	10.16	11.99	13.43	18.25	24.17	33.19		*	+	
NT2RM4002061	3.42	2.42	3.12	3.99	4.28	3.66	2.26	2.93	1.81	*	+		
NT2RM4002062	6.37	2.90	3.38	2.10	2.75	3.44	2.98	2.78	3.12				
NT2RM4002063	8.92	6.28	4.96	9.35	7.20	6.28	7.35	7.35	6.46				
NT2RM4002066	5.12	2.57	2.72	3.13	3.43	2.84	3.67	3.65	2.97				
NT2RM4002067	1.89	1.36	1.11	3.88	3.13	3.49	1.44	3.55	1.91	**	+		
NT2RM4002073	3.81	3.18	2.17	3.78	3.91	3.14	2.82	4.59	3.46				
NT2RM4002074	3.75	3.15	4.02	2.89	4.67	3.46	2.89	2.92	2.59		*	-	
NT2RM4002075	1.30	1.13	1.76	2.76	2.64	2.94	1.69	2.40	1.5	**	+		
NT2RM4002076	4.00	1.21	3.46	2.32	2.53	2.49	2.84	3.24	1.6				
NT2RM4002078	12.66	8.15	5.73	7.75	7.44	9.12	8.77	7.66	8.72				
NT2RM4002081	5.48	5.00	3.54	7.62	9.31	8.00	5.52	7.35	6.24	**	+		
NT2RM4002083	4.26	2.31	2.02	3.34	2.38	2.66	2.89	2.98	2.86				
NT2RM4002093	3.89	2.69	2.12	7.05	6.79	4.47	2.74	4.50	3.5	*	+		
NT2RM4002109	5.34	3.93	2.60	5.27	7.18	5.20	3.25	3.84	4.24				
NT2RM4002115	3.73	2.51	2.56	3.60	4.16	3.32	2.9	3.99	2.74				
NT2RM4002118	2.39	1.49	2.46	3.46	6.34	3.85	3.47	4.78	5.61		*	+	
NT2RM4002128	1.76	1.98	1.98	2.53	2.32	2.56	1.95	1.95	1.45	**	+		
NT2RM4002137	5.40	3.31	3.77	3.32	5.16	4.10	4.08	2.63	2.49				
NT2RM4002139	6.38	4.93	5.07	14.74	15.06	13.57	6.58	7.18	6.59	**	+		
NT2RM4002140	7.07	3.90	5.01	9.78	11.72	9.95	6.8	5.99	6.18	**	+		
NT2RM4002145	5.69	2.65	3.96	6.30	6.51	4.16	4.2	6.86	5.05				
NT2RM4002146	12.58	8.18	8.37	8.91	7.31	8.60	4.94	6.93	3.9				
NT2RM4002161	1.51	1.71	1.05	2.14	2.32	1.65	1.38	2.18	1.6				
NT2RM4002174	2.04	1.62	2.29	4.40	6.82	5.43	2.41	4.19	3.45	**	+		
NT2RM4002178	4.27	1.80	4.02	7.72	6.53	7.07	4.59	6.24	4.61	*	+		
NT2RM4002180	14.71	6.92	6.30	9.50	9.96	6.78	4.56	4.83	5.69				
NT2RM4002185	5.31	3.85	4.04	4.39	4.78	3.75	5.7	4.91	5.17				

73/180

【0714】

【表243】

1623

1624

NT2RM4002189	27.09	13.74	15.74	14.48	15.82	14.29	21.97	17.24	13.45				
NT2RM4002194	14.06	6.46	6.54	8.20	8.96	5.67	5.24	4.78	7.73				
NT2RM4002198	9.72	5.05	4.64	9.60	7.14	7.42	3.99	6.05	4.24				
NT2RM4002205	6.04	2.24	4.01	10.17	8.07	7.85	3.89	6.45	5.48	*	+		
NT2RM4002213	8.85	5.39	4.89	8.71	11.13	8.58	6.75	7.49	6.59				
NT2RM4002216	13.98	11.40	13.83	9.67	12.26	12.25	5.51	6.10	8.64		**	-	
NT2RM4002226	11.71	3.35	5.45	7.00	6.75	5.32	2.56	2.81	2.06				
NT2RM4002237	12.13	5.23	4.66	6.69	6.79	4.62	5.28	4.25	5.13				
NT2RM4002248	3.83	1.22	1.76	2.57	3.67	3.49	1.94	2.86	3.13				
NT2RM4002251	4.23	2.41	3.59	5.58	5.63	2.99	3.14	4.22	3.57				
NT2RM4002256	9.61	4.69	5.30	9.65	8.00	8.72	6.39	6.24	6.69				
NT2RM4002262	2.51	1.66	3.08	3.94	4.02	2.93	2.34	4.64	4.87				
NT2RM4002266	3.81	3.04	1.77	5.13	5.13	3.56	1.74	3.97	3				
NT2RM4002276	6.07	4.19	4.53	7.03	5.98	6.29	6.42	7.40	4.56				
NT2RM4002278	5.55	3.50	2.06	5.22	5.68	3.41	2.26	1.92	4.58				
NT2RM4002281	10.82	3.97	3.78	8.02	12.45	5.87	8.47	7.73	8.03				
NT2RM4002287	4.73	2.14	2.11	4.48	2.86	2.45	3.19	4.14	1.9				
NT2RM4002294	3.56	2.28	1.67	6.99	5.40	3.49	3.08	4.20	3.5				
NT2RM4002298	4.25	1.83	2.68	5.32	3.86	5.10	6.58	7.80	6.73		**	+	
NT2RM4002301	2.19	2.10	1.85	3.43	4.22	3.48	1.84	3.94	2.05	**	+		
NT2RM4002306	4.28	2.89	2.26	4.38	4.65	4.86	3.01	4.34	2.42				
NT2RM4002323	4.07	3.11	3.95	9.92	6.06	6.87	4.61	4.01	2.32	*	+		
NT2RM4002334	48.90	21.85	22.61	35.78	25.59	28.97	30.63	31.70	22.58				
NT2RM4002339	2.06	1.58	1.46	1.24	1.64	1.38	3.19	1.21	1.93				
NT2RM4002344	3.34	2.36	2.32	3.06	3.36	3.28	1.98	2.28	1.57				
NT2RM4002349	3.14	4.48	1.33	2.81	6.18	3.52	3.97	3.81	7.59				
NT2RM4002352	2.56	1.55	1.37	2.09	1.90	1.83	1.8	2.16	1.75				
NT2RM4002362	10.19	5.95	5.50	3.14	3.38	3.88	2.99	2.56	2.32		*	-	
NT2RM4002373	3.73	2.27	4.81	3.06	4.43	4.48	1.89	2.78	3.21				
NT2RM4002374	2.46	1.36	2.00	4.92	6.83	2.91	2.01	2.17	2.46				
NT2RM4002376	3.65	2.05	2.36	5.15	3.88	5.04	5.2	2.97	2.99	*	+		
NT2RM4002383	5.41	2.46	3.35	8.94	8.52	7.85	5.76	4.08	7.78	**	+		
NT2RM4002398	7.22	2.53	2.49	3.89	3.09	3.46	2.47	3.20	2.59				
NT2RM4002398	4.68	2.42	2.88	5.08	6.85	4.30	3.82	2.28	3.63				
NT2RM4002409	2.87	2.53	3.04	4.21	5.07	3.80	3.49	3.93	3.64	*	*	+	+
NT2RM4002414	5.03	1.84	3.97	3.80	4.16	6.28	4.49	4.44	4.68				
NT2RM4002438	5.21	2.42	2.20	4.07	3.59	4.94	3.44	3.46	2.5				
NT2RM4002440	4.95	2.33	3.53	5.69	5.26	3.20	3.34	4.02	4.39				
NT2RM4002446	6.41	3.72	3.77	5.16	5.23	4.99	5.81	3.91	5.57				
NT2RM4002450	7.34	5.13	5.19	4.41	3.88	3.16	3.9	3.82	4.13				
NT2RM4002452	4.76	3.56	2.63	3.31	4.00	4.75	2.58	2.59	2.32				
NT2RM4002457	3.97	2.35	2.27	5.42	4.08	5.14	4.64	3.85	2.87	*	+		
NT2RM4002458	2.05	1.17	1.07	1.55	3.27	2.46	2.27	3.06	1.92				
NT2RM4002460	1.51	0.75	1.48	0.65	1.16	0.85	1.55	1.39	1.26				
NT2RM4002464	2.69	1.95	2.48	3.72	3.71	4.31	2.38	2.92	1.83	**	+		
NT2RM4002479	6.89	5.60	6.27	9.61	8.13	4.62	4.88	6.96	5.42				
NT2RM4002482	35.61	22.59	16.97	21.71	19.69	20.16	30.11	17.88	24.23				
NT2RM4002489	15.59	8.96	10.80	10.87	12.64	11.89	10.58	8.12	11.95				
NT2RM4002492	3.66	2.45	2.96	3.64	2.32	2.09	3.63	3.17	2.29				
NT2RM4002499	39.72	27.06	27.17	54.95	67.76	43.13	21.05	20.01	15.47	*	+		
NT2RM4002504	10.06	5.00	4.83	15.16	13.66	11.30	9.77	10.12	11.17	*	+		
NT2RM4002506	3.00	2.28	3.10	3.05	3.95	4.66	3.19	3.46	3.27				
NT2RM4002510	1.71	1.62	1.42	3.05	3.64	3.86	2.57	2.61	2.07	**	+	*	+
NT2RM4002527	1.36	1.99	1.93	1.99	2.17	2.01	1.62	2.61	1.13				
NT2RM4002532	8.36	3.92	4.29	7.17	9.98	8.89	6.69	5.89	6.32				
NT2RM4002534	5.34	2.37	2.56	3.48	4.24	3.83	3.66	4.16	3.67				

74/180

【0715】

【表244】

1625

1626

NT2RM4002535	8.63	5.41	4.92	15.46	13.83	13.63	8.73	8.80	8.02	**	+		
NT2RM4002534	3.24	2.37	1.91	1.77	3.57	2.77	2.58	2.42	1.39				
NT2RM4002533	3.05	3.08	3.12	4.82	3.64	4.78	4.67	4.24	3.5	*	+	+	+
NT2RM4002565	4.27	2.27	3.74	8.08	6.46	7.43	4.53	4.18	4.36	**	+		
NT2RM4002567	2.07	1.22	2.13	2.02	9.16	3.14	2.05	3.10	2.71				
NT2RM4002571	4.37	2.84	3.54	4.69	4.81	3.75	4.16	4.57	3.27				
NT2RM4002572	6.03	2.28	2.98	6.08	6.68	4.99	9.74	8.44	9.5			*	+
NT2RM4002577	2.75	1.19	0.59	1.71	1.14	1.51	6.59	4.68	6.71	**	+		
NT2RM4002583	3.95	2.68	2.93	2.91	3.44	3.57	3.57	3.67	3.56				
NT2RM4002584	6.72	4.52	4.49	7.70	8.13	5.82	4.85	4.09	4.77				
NT2RM4002593	11.06	6.50	9.20	6.84	5.82	5.78	2.04	3.51	4.47			*	-
NT2RM4002594	4.49	2.50	2.60	5.70	6.28	5.59	4.77	7.23	6.06	*	+	+	+
NT2RM4002604	4.69	2.15	3.00	3.62	4.47	4.27	3.38	3.51	3.57				
NT2RM4002614	2.09	1.88	1.83	3.05	2.85	2.71	1.21	3.15	1.87	**	+		
NT2RM4002616	5.30	2.89	2.15	2.37	1.56	2.52	2.81	1.79	2.9				
NT2RM4002623	8.57	2.95	4.75	3.25	4.49	3.44	2.87	3.18	2.88				
NT2RM4002634	1.64	1.74	1.43	1.95	2.12	2.72	2.59	3.50	2.79			**	+
NT2RM4002636	5.12	3.99	4.07	4.89	3.26	2.51	3.1	3.30	2.24			*	-
NT2RP1000002	4.91	2.69	3.55	5.37	6.59	6.81	5.02	6.11	5.97	*	+		
NT2RP1000006	3.58	2.73	3.36	3.30	5.24	3.97	3.46	5.04	3.59				
NT2RP1000015	0.58	0.54	1.13	1.73	1.75	2.13	1.06	2.60	1.34	**	+		
NT2RP1000018	0.26	0.38	0.59	1.15	1.19	1.44	1.05	2.21	0.52	**	+		
NT2RP1000034	281.35	132.61	141.44	137.16	124.07	106.57	66.03	58.57	54.32				
NT2RP1000035	3.85	3.38	2.73	3.79	4.44	3.26	2.6	2.77	2.19				
NT2RP1000040	1.60	1.01	1.16	1.82	1.72	0.90	1.72	1.93	1.4				
NT2RP1000042	0.16	0.83	0.49	1.42	1.37	0.52	0.89	2.70	1.63				
NT2RP1000048	3.91	1.94	1.67	2.45	3.78	2.00	3.04	5.80	4.69				
NT2RP1000050	2.17	1.06	1.90	2.79	3.16	3.31	1.43	4.06	2.02	*	+		
NT2RP1000056	29.42	14.22	19.60	15.96	16.06	15.82	8.26	10.94	9.03				
NT2RP1000058	1.76	1.01	1.59	2.63	1.51	1.74	0.73	1.28	0.3				
NT2RP1000063	2.86	1.68	1.32	1.33	2.84	1.66	1.17	1.53	1.43				
NT2RP1000068	2.57	1.65	0.98	2.49	2.52	1.99	1.28	2.09	2.14				
NT2RP1000072	111.07	54.80	68.45	57.17	59.86	64.56	51.74	45.59	52.17				
NT2RP1000073	0.97	0.59	0.56	1.83	1.57	2.36	0.84	2.78	1.72	*	+		
NT2RP1000078	3.33	1.48	2.67	2.36	2.30	2.50	1.17	3.68	1.39				
NT2RP1000079	2.67	0.92	1.74	2.69	2.08	2.10	4.5	6.28	4.63			*	+
NT2RP1000080	7.28	4.50	5.28	5.11	5.46	5.42	2.3	4.02	3.44				
NT2RP1000086	4.35	3.00	3.48	3.24	3.23	2.33	1.02	2.72	1.4			*	-
NT2RP1000087	5.00	2.82	2.77	4.73	5.17	3.70	4.25	2.63	3.17				
NT2RP1000089	21.30	13.02	9.99	15.70	10.56	8.76	7.11	5.03	7.52				
NT2RP1000090	62.12	34.52	35.37	65.14	57.48	42.93	29.21	27.16	16.48				
NT2RP1000100	2.17	0.88	1.25	1.24	1.63	1.66	0.75	2.69	2.15				
NT2RP1000101	6.92	3.86	4.62	6.27	8.56	8.35	6.29	5.31	6.14				
NT2RP1000111	3.13	2.02	3.20	4.79	4.46	1.70	2.06	3.98	4.56				
NT2RP1000112	1.19	1.17	1.40	1.98	2.39	2.90	2.08	3.24	1.09	*	+		
NT2RP1000124	2.04	1.79	2.18	5.67	6.32	7.61	0.92	3.26	4.08	**	+		
NT2RP1000125	13.33	6.69	5.55	16.93	13.49	11.53	18.17	14.66	19.62			*	+
NT2RP1000129	8.42	3.01	2.92	5.33	4.43	3.32	3.8	3.24	4.62				
NT2RP1000130	3.80	3.59	3.16	6.14	5.63	6.01	3.49	3.06	4.37	**	+		
NT2RP1000134	2.77	1.66	1.73	4.97	6.35	4.78	3.19	4.61	2.92	**	+		
NT2RP1000163	2.54	1.56	0.69	1.65	3.20	1.85	0.24	3.07	0.88				
NT2RP1000170	1.25	0.62	0.44	1.83	1.94	1.90	0.89	3.09	1.57	**	+		
NT2RP1000174	0.77	0.39	0.59	0.80	1.14	0.73	0.83	1.30	0.25				
NT2RP1000181	15.66	7.51	13.59	20.37	20.72	18.84	8.95	8.78	5.68	*	+		
NT2RP1000191	2.05	1.96	1.05	3.54	1.96	2.31	1.34	1.45	2.86				
NT2RP1000202	1.43	1.24	0.92	2.91	2.20	1.99	0.8	2.37	2.35	*	+		

75/180

[0716]

[表245]

1627

1628

NT2RP1000239	0.54	0.73	0.33	1.02	1.34	0.58	0.16	1.81	1.18				
NT2RP1000243	0.84	0.90	0.58	2.06	1.73	0.89	0.86	2.41	1.44				
NT2RP1000255	0.75	0.34	1.01	1.49	0.71	0.89	0.75	1.85	0.92				
NT2RP1000259	1.78	1.74	1.10	4.78	3.63	3.84	2.82	4.39	2.57	**	+	+	+
NT2RP1000261	1.08	0.77	0.32	2.74	1.80	1.60	0.7	2.51	1.42	+			
NT2RP1000269	12.70	6.05	5.79	12.05	12.78	10.09	7.5	10.31	8.4				
NT2RP1000271	65.05	27.46	27.30	118.92	88.05	70.43	44.58	28.04	22.55				
NT2RP1000272	15.64	8.87	8.62	11.91	10.97	10.04	8.77	5.04	6.08				
NT2RP1000279	3.64	2.60	2.52	4.01	4.52	4.50	3.4	3.60	2.95	+			
NT2RP1000290	31.80	25.40	25.39	36.52	40.12	40.15	26.39	22.99	29.24	**	+		
NT2RP1000283	8.90	5.15	6.17	9.07	11.34	10.12	7.62	7.73	8.67				
NT2RP1000300	21.75	19.20	18.07	20.53	28.21	20.72	16.45	24.53	12.12				
NT2RP1000324	12.47	5.32	8.89	10.68	13.57	9.75	6.98	9.83	9.18				
NT2RP1000325	91.19	35.26	49.60	54.44	61.67	55.26	47.32	30.15	44.99				
NT2RP1000326	10.60	7.28	6.00	12.46	8.25	10.43	7.71	8.51	5.43				
NT2RP1000331	13.85	7.24	6.82	12.25	10.31	7.00	5.01	4.72	3.71				
NT2RP1000333	12.54	6.22	6.02	8.86	8.17	8.74	6.53	7.71	7.88				
NT2RP1000336	1.87	1.73	1.02	1.35	1.53	1.21	3.14	2.70	2.83			+	+
NT2RP1000347	2.75	2.10	2.68	2.09	2.48	2.62	1.53	2.25	0.84				
NT2RP1000348	1.47	0.48	0.33	1.45	1.42	1.72	1.13	1.89	0.66				
NT2RP1000349	0.93	0.52	0.64	1.41	1.77	1.72	0.95	0.90	1.19	**	+		
NT2RP1000353	40.50	18.12	20.02	27.21	16.43	19.17	10.71	8.40	12.57				
NT2RP1000356	39.98	22.39	20.90	32.15	26.26	25.06	14.83	10.10	14.28				
NT2RP1000357	13.61	7.81	6.20	11.20	13.90	12.68	8.98	8.00	11.38				
NT2RP1000358	11.64	5.39	5.27	10.20	9.77	8.75	7.77	6.88	9.19				
NT2RP1000360	26.32	15.93	17.17	17.83	19.58	19.99	16.48	15.94	15.67				
NT2RP1000363	22.05	14.66	16.07	21.39	24.54	24.53	22.26	17.18	17.26				
NT2RP1000376	5.84	3.91	5.30	4.51	6.40	6.42	7.18	6.13	5.77				
NT2RP1000386	31.79	21.04	23.39	64.26	64.31	34.90	56.81	60.95	58.22	+	**	+	+
NT2RP1000487	0.29	0.73	0.45	0.62	0.61	0.29	1.08	0.88	0.22				
NT2RP1000489	2.22	1.91	0.68	2.83	3.38	2.80	2.71	1.86	1.7				
NT2RP1000413	7.71	3.51	3.65	7.04	7.63	7.01	5.32	4.65	6.75				
NT2RP1000416	2.07	0.73	0.71	1.73	2.70	2.64	1.38	1.53	1.42				
NT2RP1000418	0.88	0.78	0.91	2.07	1.77	2.03	1.84	2.71	1.4	**	+	+	+
NT2RP1000420	0.51	0.68	0.34	1.31	0.46	1.21	1.33	1.51	1.45				
NT2RP1000434	0.66	0.29	2.53	1.80	1.28	1.15	1.63	2.36	0.97				
NT2RP1000439	13.59	10.41	10.76	8.22	11.89	8.15	6.48	6.20	3.53			+	+
NT2RP1000443	1.67	1.60	1.02	3.09	3.95	2.04	3.35	1.76	1.48				
NT2RP1000447	2.13	0.82	0.90	2.07	1.85	1.21	1.39	1.67	1.12				
NT2RP1000448	1.39	0.47	0.72	0.68	1.75	1.34	1.82	1.77	0.69				
NT2RP1000451	5.40	2.45	1.97	5.69	5.15	3.49	1.66	2.36	1.96				
NT2RP1000458	22.07	12.50	14.79	20.35	29.47	24.03	21.83	19.22	26.03				
NT2RP1000466	19.74	9.97	12.40	17.61	20.40	21.09	17.72	15.83	18.24				
NT2RP1000468	14.77	10.71	12.70	18.32	19.61	21.10	14.71	11.30	11.86	**	+		
NT2RP1000468	3.47	2.54	4.12	7.07	8.07	7.42	3.93	5.61	4.57	**	+		
NT2RP1000479	14.45	6.40	6.23	5.28	6.94	7.41	8.62	6.71	6.97				
NT2RP1000477	0.33	0.76	0.21	0.93	1.49	0.73	0.8	1.04	0.52				
NT2RP1000478	2.01	1.44	1.12	1.74	1.18	2.18	1.98	3.01	1.97				
NT2RP1000481	3.26	1.45	1.19	1.27	1.08	1.24	0.92	2.02	0.85				
NT2RP1000493	1.13	0.65	0.54	1.16	1.49	1.41	1.57	2.12	0.89	+			
NT2RP1000513	8.57	3.43	5.13	11.73	10.43	8.69	10.51	9.55	9.33				
NT2RP1000532	9.74	3.47	5.93	6.13	9.61	9.77	8.53	8.00	7.9				
NT2RP1000533	2.49	0.79	1.92	2.45	2.66	3.02	1.21	2.77	1.5				
NT2RP1000544	2.42	0.99	0.69	2.39	1.44	1.14	1.43	1.13	2.11				
NT2RP1000547	0.17	0.54	0.23	0.77	0.69	0.77	0.43	1.67	0.73	+			
NT2RP1000553	1.62	1.44	0.64	0.50	0.71	0.60	1.24	2.56	1.59				

1629

1630

NT2RF1000567	1.21	0.33	0.63	1.21	1.30	2.41	2.12	3.42	1.77		*	+
NT2RF1000574	1.82	0.32	0.03	23.76	28.12	20.34	4.23	4.69	3.79	**	+	+
NT2RF1000577	1.22	0.49	0.73	1.46	1.85	1.75	1.18	2.92	1.35	*	+	
NT2RF1000579	0.79	0.65	0.57	1.33	1.34	1.32	1.35	2.50	0.76	**	+	
NT2RF1000581	1.36	0.66	1.82	2.04	1.53	1.78	1.95	2.51	1.09			
NT2RF1000593	2.64	0.66	1.75	2.65	2.96	1.71	1.41	0.83	1.4			
NT2RF1000604	11.50	7.94	7.40	3.94	3.98	3.21	2.12	2.31	2.08	*	**	-
NT2RF1000609	2.53	2.00	0.54	1.02	1.56	1.09	1.82	2.61	1.48			
NT2RF1000613	1.94	0.88	0.65	1.32	0.99	1.16	0.85	2.58	1.01			
NT2RF1000622	1.32	0.92	0.99	1.13	1.63	1.80	1.57	3.98	2.19			
NT2RF1000627	5.47	2.19	3.87	5.94	4.15	4.81	4.23	6.27	4.91			
NT2RF1000629	1.49	0.86	0.85	1.86	1.84	2.88	2.18	2.88	1.87		*	+
NT2RF1000630	5.89	2.85	5.42	13.99	11.47	13.46	7.36	6.55	7.16	**	+	
NT2RF1000639	2.68	1.18	0.53	1.84	1.97	0.94	1.56	1.83	1.5			
NT2RF1000640	81.74	37.60	35.82	57.27	52.32	39.58	48.18	42.34	41.38			
NT2RF1000646	7.82	4.91	3.97	8.29	9.40	9.31	5.5	5.31	6.52	*	+	
NT2RF1000659	6.71	2.34	3.90	4.05	6.32	6.12	3.31	4.60	4.15			
NT2RF1000674	4.71	2.08	3.93	3.76	7.16	7.25	3.17	4.95	4.5	*	+	
NT2RF1000677	9.51	6.01	6.41	8.66	8.51	8.83	7.33	7.01	8.68			
NT2RF1000679	1.23	0.42	0.82	1.73	1.38	1.63	1.09	2.06	0.76	*	+	
NT2RF1000688	4.67	2.07	2.03	5.85	5.34	3.72	3.1	4.12	2.68			
NT2RF1000689	2.83	0.64	1.04	1.11	1.67	0.84	1.37	0.88	0.83			
NT2RF1000695	1.62	1.12	1.10	1.18	2.99	1.24	1	0.87	0.88			
NT2RF1000701	0.90	0.82	0.62	0.83	0.27	1.25	0.87	1.19	1.4			
NT2RF1000703	0.76	0.35	1.53	0.66	1.47	1.82	0.6	1.47	2.57			
NT2RF1000713	0.23	0.42	0.37	0.34	0.89	0.44	0.17	1.44	1.47			
NT2RF1000721	10.57	6.36	5.67	7.28	13.00	9.92	8.49	9.05	8.17			
NT2RF1000730	2.55	1.65	1.97	4.38	3.90	3.35	1.75	3.65	2.95	*	+	
NT2RF1000733	4.46	2.99	3.71	5.44	5.04	3.14	1.44	3.93	4.16			
NT2RF1000738	28.84	10.50	11.79	17.48	18.85	18.44	15.99	11.63	12.72			
NT2RF1000739	14.40	7.16	8.58	10.60	12.85	8.63	11.15	9.94	11.2			
NT2RF1000740	3.66	1.37	2.15	2.84	4.09	2.86	2.91	2.60	3.23			
NT2RF1000746	1.31	0.65	0.82	1.32	1.26	0.89	1.26	2.13	2.46			
NT2RF1000750	9.51	4.76	5.09	7.09	6.45	6.48	4.95	5.43	4.72			
NT2RF1000751	77.49	46.65	53.99	41.34	32.45	28.11	17.67	20.76	21.6	*	-	
NT2RF1000767	1.53	0.63	1.06	1.68	1.34	1.25	1.21	2.74	2.71			
NT2RF1000769	4.65	2.64	3.84	2.57	3.18	2.72	4.13	4.77	3.22			
NT2RF1000789	1.51	0.92	0.80	2.30	1.18	0.64	1.37	0.96	0.77			
NT2RF1000782	5.21	2.12	2.72	11.13	10.26	10.71	6.05	7.66	6.54	**	+	+
NT2RF1000796	6.49	4.06	3.11	4.93	5.23	3.73	4.82	3.98	7.13			
NT2RF1000797	11.72	5.77	5.28	6.51	8.45	5.34	7.81	7.98	9.33			
NT2RF1000800	0.13	0.54	1.00	1.07	2.16	1.97	0.82	3.18	1.42	*	+	
NT2RF1000825	3.33	1.37	1.55	2.64	2.23	1.50	1.34	2.32	1.31			
NT2RF1000833	6.35	2.68	2.53	4.24	4.98	4.14	2.29	4.43	2.42			
NT2RF1000834	16.60	5.93	7.79	8.68	7.93	6.33	6.47	7.84	5.03			
NT2RF1000836	1.43	1.06	0.85	1.19	1.20	0.59	2.19	1.50	0.63			
NT2RF1000837	6.20	2.33	2.35	4.62	5.53	5.38	4.6	3.52	3.49			
NT2RF1000846	1.21	0.89	0.89	1.89	2.60	1.73	1.96	1.80	1.08	*	+	
NT2RF1000847	2.27	1.79	1.06	1.99	2.12	2.09	2.78	1.80	2.3			
NT2RF1000851	10.08	6.27	7.87	9.89	12.49	7.13	7.78	9.66	7.43			
NT2RF1000856	9.90	5.85	7.31	20.58	23.87	20.13	15.75	15.89	19.71	**	+	+
NT2RF1000860	7.91	5.43	8.96	10.11	6.72	7.04	5.54	7.17	4.85			
NT2RF1000902	2.64	0.85	0.61	5.04	4.02	3.81	3.86	2.25	3.35	*	+	
NT2RF1000903	7.75	3.79	2.92	4.96	6.61	5.49	5.15	5.13	5.52			
NT2RF1000905	3.44	2.09	1.19	3.49	2.21	2.49	3.41	1.72	2.10			
NT2RF1000915	15.16	7.68	7.64	8.98	6.57	7.27	3.44	4.20	4			

77/180

【0718】

【表247】

1631

1632

NZTRP1000916	3.20	2.97	0.92	3.11	2.12	2.20	3.16	2.17	2.19								
NZTRP1000921	1.84	1.45	1.78	3.53	2.23	2.78	2.9	2.99	2.53	+	+	+	+	+	+	+	+
NZTRP1000943	1.83	0.78	1.29	5.94	5.07	3.31	7.05	7.60	6.55	++	+	+	+	+	+	+	+
NZTRP1000944	3.54	2.52	3.09	5.21	4.55	4.65	2.55	2.42	2.55	++	+	+	+	+	+	+	+
NZTRP1000947	6.89	4.11	3.31	6.97	6.41	5.03	5.81	4.15	4.54								
NZTRP1000954	5.12	2.35	2.15	5.92	4.95	4.84	4.75	3.63	4.18								
NZTRP1000958	20.62	10.44	1.43	11.21	10.24	6.49	7.05	5.48	7.18								
NZTRP1000959	72.36	35.16	43.30	53.44	48.85	40.35	20.64	19.16	22.61								
NZTRP1000966	36.86	19.10	21.19	22.56	35.39	24.14	15.07	9.91	18.23								
NZTRP1000974	10.91	8.14	8.28	18.92	22.10	19.21	14.69	15.24	13.39	++	+	+	+	+	+	+	+
NZTRP1000980	3.63	2.59	2.91	3.75	4.02	3.96	2.97	3.22	2.22								
NZTRP1000981	4.96	3.42	4.61	4.59	5.02	3.62	2.94	3.11	2.77								
NZTRP1000988	2.69	1.97	1.73	4.25	5.22	4.19	3.95	3.30	3.66	++	+	+	+	+	+	+	+
NZTRP1001002	6.75	4.73	2.89	3.13	4.46	2.79	4.86	5.58	5.21								
NZTRP1001004	1.76	1.26	0.75	1.72	1.80	2.22	3.2	2.14	2.89								
NZTRP1001007	1.72	0.91	0.86	2.02	1.84	1.75	3	2.58	3.22								
NZTRP1001011	4.98	3.03	2.17	7.06	8.67	6.46	5.23	4.65	5.76	++	+	+	+	+	+	+	+
NZTRP1001013	3.60	3.50	3.48	9.46	12.09	7.99	6.88	5.63	8.02	++	+	+	+	+	+	+	+
NZTRP1001014	3.96	3.16	3.28	4.93	3.71	4.01	3.71	3.05	2.43								
NZTRP1001020	3.23	1.24	1.06	2.23	1.86	1.47	2.29	2.09	1.68								
NZTRP1001023	261.06	118.84	124.95	113.92	104.93	83.66	236.2	219.46	213.5								
NZTRP1001037	12.10	6.08	4.74	9.03	7.91	6.47	4.01	4.15	4.04								
NZTRP1001031	2.17	1.05	0.67	1.79	1.31	1.73	0.62	1.86	1.93								
NZTRP1001033	2.89	1.62	1.96	3.31	4.49	3.57	2.4	3.46	2.46	+	+	+	+	+	+	+	+
NZTRP1001043	2.56	1.34	2.04	5.44	5.57	4.27	4.68	4.47	4.7	++	+	+	+	+	+	+	+
NZTRP1001045	55.87	37.46	39.12	31.66	32.21	26.52	26.73	25.41	27.84								
NZTRP1001073	18.17	10.94	13.65	7.43	11.62	10.45	3.3	5.82	3.6								
NZTRP1001079	6.27	4.29	4.83	7.17	5.68	5.81	5.84	5.09	3.64								
NZTRP1001080	4.59	3.36	2.02	3.32	2.67	3.66	3.81	3.01	2.62								
NZTRP1001113	2.09	1.06	0.43	0.85	1.89	1.25	1.74	2.63	1.22								
NZTRP1001159	22.23	15.34	13.51	27.36	29.04	20.75	11.14	12.23	9.12								
NZTRP1001173	2.37	0.91	1.48	10.20	7.72	8.04	6.93	5.00	6.33	++	+	+	+	+	+	+	+
NZTRP1001176	5.14	3.86	5.35	6.46	6.12	5.31	4.46	5.39	4.12								
NZTRP1001177	3.79	2.64	3.45	7.23	6.84	5.24	5.18	4.11	3.16	+	+	+	+	+	+	+	+
NZTRP1001188	4.77	2.20	2.83	10.28	7.74	6.42	4.72	4.39	3.75	+	+	+	+	+	+	+	+
NZTRP1001199	2.06	1.25	1.14	4.62	4.88	3.76	2.05	2.71	1.7	++	+	+	+	+	+	+	+
NZTRP1001205	19.37	11.82	11.58	17.19	17.16	12.69	6.66	6.05	4.62								
NZTRP1001215	5.66	2.61	2.14	2.79	3.86	3.71	2.65	3.10	2.8								
NZTRP1001225	5.42	2.06	1.65	2.88	2.39	2.40	3.21	4.49	4.21								
NZTRP1001245	3.12	2.63	4.04	4.32	4.51	4.91	3.1	5.42	4.42	+	+	+	+	+	+	+	+
NZTRP1001247	1.41	0.44	0.55	0.62	0.90	1.10	0.75	2.81	1								
NZTRP1001248	2.58	2.07	1.62	3.98	2.41	2.41	1.39	3.80	1.81								
NZTRP1001253	6.69	3.25	3.71	6.33	4.35	5.83	4.57	5.25	3.74								
NZTRP1001286	3.18	1.26	2.31	4.52	3.67	4.87	3.96	3.81	2.61	+	+	+	+	+	+	+	+
NZTRP1001294	9.78	2.41	4.54	3.50	4.67	2.27	2.68	1.87	2.59								
NZTRP1001302	8.57	3.22	3.02	3.18	3.24	3.37	2.74	2.23	2.95								
NZTRP1001310	9.73	5.23	5.10	9.63	10.00	7.15	7.46	7.70	6.61								
NZTRP1001311	18.47	7.91	7.87	3.75	8.43	7.25	3.98	5.42	3.54								
NZTRP1001313	10.94	5.16	4.72	12.65	11.32	9.22	3.47	5.55	4.61								
NZTRP1001324	3.38	2.26	1.54	3.44	2.03	2.97	2.03	3.34	1.99								
NZTRP1001349	3.51	1.77	2.13	2.29	2.35	2.91	2.76	4.09	1.76								
NZTRP1001361	9.53	5.57	12.07	15.75	14.43	10.15	3.96	7.68	5.72								
NZTRP1001379	9.49	3.63	4.16	6.43	5.54	3.66	4.65	4.16	4.18								
NZTRP1001385	6.18	2.32	2.60	4.81	6.35	3.73	2.76	3.62	3.67								
NZTRP1001395	5.45	2.82	3.04	4.04	3.63	2.71	4.99	4.64	3.44								
NZTRP1001410	18.25	3.37	10.42	15.62	9.58	11.66	11.21	9.39	10.03								

1633

1634

NT2RF1001424	2.87	1.62	0.72	3.11	2.58	2.58	1.61	3.38	2.21										
NT2RF1001432	2.47	1.17	2.41	2.23	2.48	1.53	1.78	3.14	1.45										
NT2RF1001449	7.62	4.22	5.10	9.69	11.61	8.75	6.99	5.82	6.74	*	+								
NT2RF1001457	4.04	2.37	2.71	3.08	3.14	2.75	2.72	2.61	3.14										
NT2RF1001459	10.76	3.49	3.82	8.95	9.17	5.61	7.87	6.73	6.96										
NT2RF1001466	22.82	9.71	11.08	9.67	7.98	7.40	7.72	5.26	6.18										
NT2RF1001475	6.67	4.07	4.28	8.53	10.26	8.11	4.73	4.80	4.35	*	+								
NT2RF1001482	11.57	4.98	6.24	6.89	3.62	4.62	2.44	2.41	2.61										
NT2RF1001494	1.38	1.05	0.95	2.03	1.52	1.37	0.9	2.19	2.18										
NT2RF1001580	2.19	2.12	1.80	1.11	1.95	1.39	1	2.88	1.81										
NT2RF1001517	1.81	0.96	1.45	2.37	1.81	2.59	1.22	2.90	1.19										
NT2RF1001540	5.66	2.57	3.71	5.28	5.66	5.56	4.29	5.21	3.47										
NT2RF1001543	8.78	3.57	3.35	10.80	11.59	6.71	5.01	4.67	5.34										
NT2RF1001546	21.79	10.60	8.72	53.53	51.38	41.78	29.72	27.77	37.59	**	+	*							
NT2RF1001550	9.54	5.59	4.56	11.19	14.45	11.40	5.56	6.79	7.13	*	+								
NT2RF1001553	6.39	3.38	2.69	4.45	3.49	2.74	3.6	4.16	2.78										
NT2RF1001555	9.92	5.57	6.23	12.43	10.74	10.45	11.69	18.08	11.64										
NT2RF1001563	4.37	1.97	2.43	3.66	4.03	3.10	1.84	4.08	2.22										
NT2RF1001569	5.25	3.17	2.27	4.32	4.47	4.21	3.54	6.70	3.89										
NT2RF1001584	8.28	4.33	4.71	6.70	8.09	6.25	5.94	7.17	6.75										
NT2RF1001599	7.22	2.05	1.29	32.60	27.43	19.18	6.56	7.36	8.71	**	+								
NT2RF1001616	3.29	0.83	1.26	2.03	2.10	1.09	2.49	2.44	3.45										
NT2RF1001654	19.86	5.14	4.62	10.80	10.51	8.45	6.66	9.40	9.83										
NT2RF1001665	1.29	1.28	0.35	1.08	1.87	1.90	0.74	2.81	0.7										
NT2RF1001679	87.88	43.02	42.15	72.20	73.59	55.81	28.48	41.49	35.04										
NT2RF1001681	21.69	14.86	18.60	13.78	17.18	10.98	7.11	14.27	11.13										
NT2RF1001694	8.51	6.03	4.96	4.21	4.41	2.94	5.31	11.65	6.79										
NT2RF2000001	6.32	1.40	2.79	3.24	2.80	2.62	3.54	4.14	4.06										
NT2RF2000006	2.04	1.48	0.96	4.50	2.92	2.33	2.69	2.50	1.61										
NT2RF2000007	10.09	4.44	5.04	3.97	3.31	4.03	3.55	1.69	1.81										
NT2RF2000008	10.88	5.03	5.27	12.65	14.30	9.35	7.5	5.73	4.32										
NT2RF2000010	1.99	1.02	0.52	2.09	3.06	2.49	2.1	2.41	2.6										
NT2RF2000011	7.02	4.29	5.02	10.56	10.46	8.08	6.55	5.23	6.43	*	+								
NT2RF2000027	3.12	1.85	1.41	5.78	3.32	2.95	2.99	2.41	1.39										
NT2RF2000028	2.89	1.81	1.90	3.51	2.63	2.74	4.34	5.32	5.48			**	+						
NT2RF2000033	1.94	1.20	2.03	2.85	3.78	4.04	1.05	2.10	0.96	*	+								
NT2RF2000040	37.68	15.23	16.54	19.89	18.06	16.95	22.42	19.65	16.11										
NT2RF2000043	9.28	3.40	4.33	7.54	7.04	6.30	5.89	6.48	6.12										
NT2RF2000045	10.41	4.33	5.29	6.44	6.23	7.07	5.45	5.93	4.14										
NT2RF2000051	12.68	6.63	7.07	5.35	6.94	5.58	5.26	5.53	4.86										
NT2RF2000054	5.27	3.29	2.87	3.88	5.04	4.42	5.28	3.65	4.48										
NT2RF2000056	4.49	2.47	2.46	3.36	3.01	3.82	3.5	3.62	3.48										
NT2RF2000057	52.52	38.64	47.28	59.49	56.29	50.39	23.72	29.39	31.7	*									
NT2RF2000067	3.42	1.83	2.49	4.64	3.08	3.41	1.5	3.38	2.02										
NT2RF2000079	8.99	4.22	3.23	5.71	5.95	7.00	8.23	3.07	7.09										
NT2RF2000076	2.83	1.15	1.15	1.86	1.61	1.73	2.7	1.97	2.2										
NT2RF2000077	10.69	4.72	3.55	9.58	8.73	8.11	7.3	4.40	8.28										
NT2RF2000079	4.88	3.21	3.11	8.07	7.12	7.59	4.5	3.56	4.48	**	+								
NT2RF2000088	3.67	3.74	2.96	4.10	4.22	2.91	4.51	4.30	4.17										
NT2RF2000091	3.05	2.14	3.45	10.95	9.06	8.83	4.37	6.05	6.43	**	+	*							
NT2RF2000092	10.83	5.23	7.63	16.92	17.59	12.32	8.09	11.12	9.14	*	+								
NT2RF2000097	2.33	2.76	2.63	4.90	4.82	3.90	2.22	2.43	2.99	**	+								
NT2RF2000098	10.38	5.79	6.50	5.56	4.26	4.65	2.67	1.61	2.03			*							
NT2RF2000108	9.83	5.39	6.38	12.17	15.62	9.37	8.01	6.04	4.82										
NT2RF2000114	2.05	1.50	1.13	3.20	1.92	2.20	3.45	2.13	2.56										
NT2RF2000116	5.05	3.16	5.23	7.97	9.36	8.63	7.01	7.36	8.27	**	+	*	+						

79/180

【0720】

【表249】

1639

1640

NT2RP2000777	29.85	20.42	21.91	16.22	17.42	15.02	12.37	12.07	13.5		*	-
NT2RP2000786	8.23	5.22	4.46	10.55	9.74	7.80	11.68	12.09	10.59		*	+
NT2RP2000793	14.01	7.42	10.26	12.19	18.41	17.74	12.21	12.81	15.75			
NT2RP2000796	6.25	2.57	4.14	5.05	5.14	3.85	3.27	4.71	3.04			
NT2RP2000809	7.70	5.02	4.14	9.32	10.55	8.44	6.87	4.85	6.03	*	+	
NT2RP2000812	6.41	3.65	3.75	7.05	6.14	5.86	4.83	5.04	3.67			
NT2RP2000814	2.40	1.13	1.50	2.03	1.96	1.78	0.9	2.54	1.09			
NT2RP2000816	5.89	1.17	2.01	3.48	3.06	4.82	3.84	4.28	3.29			
NT2RP2000818	2.61	0.71	0.86	3.13	3.87	2.75	2.08	1.63	3.25			
NT2RP2000819	2.57	1.24	1.34	1.88	1.49	1.77	1.81	2.05	1.32			
NT2RP2000841	2.46	0.72	1.21	2.94	1.98	3.02	1.06	2.75	1.48			
NT2RP2000842	1.34	0.54	1.09	1.95	1.43	1.71	2.84	2.70	1.53	*	+	
NT2RP2000845	12.78	5.61	3.57	11.56	12.23	11.13	7.34	7.10	8.72			
NT2RP2000853	2.24	1.48	1.52	2.02	1.72	1.96	1.61	2.25	1.68			
NT2RP2000880	10.87	4.76	7.03	10.28	10.84	10.60	7.87	8.04	7.97			
NT2RP2000892	3.07	1.45	2.10	2.15	3.52	2.03	2.6	3.34	2.68			
NT2RP2000894	2.45	1.27	1.87	2.80	3.03	2.60	3.77	5.13	5.17		**	+
NT2RP2000903	2.42	1.74	2.17	15.91	10.43	12.06	3.76	4.80	3.91	**	+	**
NT2RP2000906	2.89	1.95	2.70	4.14	5.17	4.16	3.32	2.67	4.12	*	+	
NT2RP2000910	2.79	1.53	2.66	6.17	5.30	4.67	3.71	4.07	3.28	**	+	*
NT2RP2000931	32.13	11.92	13.53	39.97	39.93	28.59	17.58	15.27	16.3			
NT2RP2000932	4.21	2.31	2.05	7.96	6.87	4.87	4.36	3.76	4.67	*	+	
NT2RP2000938	19.54	10.59	13.57	13.71	16.06	13.76	9.46	10.81	12.03			
NT2RP2000943	4.61	2.00	2.25	2.99	4.17	3.48	6.66	6.59	6.2	*	+	
NT2RP2000957	2.25	1.38	1.92	2.45	2.33	2.46	1.28	3.48	2.23			
NT2RP2000958	6.62	2.75	4.11	5.71	4.71	5.65	4.44	6.65	3.45			
NT2RP2000959	5.43	1.74	2.79	6.81	7.31	5.96	7.7	6.58	8.28	*	+	*
NT2RP2000965	8.62	7.11	7.91	6.90	6.39	7.29	4.61	4.19	4.83		**	-
NT2RP2000970	6.70	2.82	2.67	8.85	8.32	8.60	5.68	4.48	4.57	*	+	
NT2RP2000973	3.87	3.35	2.21	3.68	3.61	1.94	3.33	3.24	2.43			
NT2RP2000985	4.15	2.39	2.33	2.87	4.28	3.35	2.71	2.53	3.95			
NT2RP2000987	2.36	1.40	1.29	2.94	3.30	3.87	2.43	3.02	3.28	*	+	*
NT2RP2000997	3.92	3.46	2.91	6.76	6.13	8.29	6.06	7.63	6.82	**	+	**
NT2RP2001004	3.02	2.00	2.80	4.39	4.00	3.80	2.57	2.72	3.03	*	+	
NT2RP2001028	1.53	1.61	1.49	3.31	2.89	2.16	1.09	3.10	1.56	*	+	
NT2RP2001036	8.99	5.09	6.28	14.47	12.09	13.66	6.21	7.37	8.86	**	+	
NT2RP2001039	2.38	1.24	0.84	2.83	2.64	1.64	1.83	1.41	1.82			
NT2RP2001044	3.60	1.75	2.33	3.81	3.95	2.60	1.92	3.42	3.51			
NT2RP2001056	8.76	6.20	3.80	10.38	10.96	8.29	5.85	5.19	6.9			
NT2RP2001065	11.06	6.53	6.66	6.07	7.52	5.67	4.84	4.18	3.98			
NT2RP2001067	3.97	2.56	1.95	4.29	2.72	3.44	1.28	3.38	2.55			
NT2RP2001070	6.27	3.18	2.94	8.92	8.75	6.08	5.11	6.42	3.18			
NT2RP2001081	7.29	3.39	2.85	9.20	10.42	10.02	6.26	8.11	6.41	*	+	
NT2RP2001087	2.47	2.17	1.24	3.46	5.06	3.87	2.98	3.13	3.05	*	+	*
NT2RP2001094	0.61	0.13	0.10	1.14	0.70	0.35	0.83	0.88	1.21	*	+	*
NT2RP2001119	6.84	4.46	3.47	7.70	9.69	7.83	4.19	5.13	8.84	*	+	
NT2RP2001127	5.97	3.17	2.14	8.14	7.01	6.94	3.37	5.51	5.47	*	+	
NT2RP2001133	6.80	4.14	3.76	7.22	8.84	6.01	3.82	6.62	4.59			
NT2RP2001137	4.85	2.38	2.65	2.75	3.98	3.93	2.74	5.27	3.23			
NT2RP2001143	3.86	1.91	2.02	3.11	3.09	2.46	1.97	4.83	1.3			
NT2RP2001149	4.02	1.34	2.11	3.88	2.95	3.29	1.85	2.88	2.33			
NT2RP2001168	13.95	5.65	7.80	16.05	15.12	13.54	11.11	11.37	10.13			
NT2RP2001173	2.96	1.32	1.35	7.72	6.56	4.53	4.19	3.26	2.72	*	+	
NT2RP2001174	4.49	3.17	1.74	5.69	5.38	5.09	5.65	3.56	3.21			
NT2RP2001184	7.71	4.21	4.96	7.15	6.32	5.98	5.09	5.61	5.63			
NT2RP2001196	1.68	0.99	1.05	1.56	1.51	1.49	1.6	1.79	2.14			

82/180

1641

1642

NT2RP2001209	3.43	3.44	2.46	6.53	4.88	4.21	3.59	2.77	3.29										
NT2RP2001218	3.11	1.72	2.13	3.51	3.65	3.23	2.31	2.98	3.88										
NT2RP2001223	5.06	2.55	3.61	3.72	4.59	2.27	3.19	3.20	3.06										
NT2RP2001226	12.72	7.29	8.85	12.01	9.47	7.65	11.46	8.46	11.8										
NT2RP2001227	6.22	4.18	3.44	6.26	5.08	5.75	7.03	4.88	5.64										
NT2RP2001232	7.29	3.90	3.93	7.87	8.17	8.48	7.29	5.90	4.44										
NT2RP2001233	14.76	8.17	8.10	14.08	19.00	21.01	13.52	10.12	10.65										
NT2RP2001245	3.69	2.29	2.63	3.56	3.59	3.28	3.42	3.62	4.39										
NT2RP2001246	2.35	0.80	3.09	3.34	4.44	4.13	4.38	7.67	6.87										
NT2RP2001248	5.55	3.73	6.74	8.43	9.77	9.29	5.65	6.17	7.45	*									
NT2RP2001278	14.16	9.13	9.94	14.63	14.49	8.30	11.4	14.47	14.26										
NT2RP2001276	2.24	1.82	0.94	3.36	2.75	2.46	3.31	2.32	2.92										
NT2RP2001277	3.77	1.80	1.15	7.12	6.46	6.90	6.6	4.91	5.92	**									
NT2RP2001279	3.82	2.12	2.26	5.58	9.49	5.69	6.49	4.63	4.63	*									
NT2RP2001295	3.75	1.96	2.66	4.93	5.60	3.83	3.62	3.11	3.56										
NT2RP2001297	104.94	62.95	78.61	112.57	111.95	109.12	28.51	42.30	59.76										
NT2RP2001301	6.22	5.96	7.50	7.48	6.39	7.90	5.94	7.38	6.32										
NT2RP2001312	16.14	10.26	15.91	20.56	19.30	16.72	18.23	19.30	23.86										
NT2RP2001327	8.14	6.35	5.95	3.76	7.30	7.36	7.73	8.61	9.09										
NT2RP2001328	18.42	9.64	9.66	24.64	22.08	22.34	13.94	10.86	12.67	*									
NT2RP2001341	17.63	7.30	6.72	12.36	9.62	10.30	8.25	8.97	14.63										
NT2RP2001347	17.63	11.15	9.87	16.21	14.33	12.17	10.57	9.73	12.31										
NT2RP2001366	10.12	8.31	6.45	18.92	23.58	18.36	11.75	11.32	14.59	**									
NT2RP2001378	8.29	6.95	6.58	6.49	8.22	6.02	7.98	9.16	9.41										
NT2RP2001381	4.07	2.97	3.94	2.90	3.52	4.42	2.95	2.69	2.85										
NT2RP2001388	3.41	3.63	3.35	6.25	9.01	7.41	5.95	6.27	6.62	**									
NT2RP2001391	210.40	161.64	144.04	393.09	492.35	288.04	175.7	224.46	230.6	*									
NT2RP2001392	7.04	3.01	3.58	4.59	5.33	4.71	8.14	5.70	5.27										
NT2RP2001394	9.60	6.22	4.32	15.24	15.30	14.78	8	5.76	7.4	**									
NT2RP2001397	15.57	11.63	10.83	8.23	11.47	9.12	4.18	3.62	3.82										
NT2RP2001408	2.42	2.39	2.33	4.87	6.19	6.06	7.4	8.87	13.18	**									
NT2RP2001409	5.20	3.88	3.54	7.39	10.57	7.94	7.53	7.30	6.48	*									
NT2RP2001428	4.15	2.99	3.26	8.92	7.75	7.19	4.98	4.32	3.55	**									
NT2RP2001433	3.65	2.45	3.55	6.47	6.38	4.42	6.23	5.04	5.49	*									
NT2RP2001427	4.90	3.28	3.58	5.81	6.42	5.73	4.13	4.89	4.51	*									
NT2RP2001438	4.31	2.09	2.32	7.25	7.90	5.77	3.53	5.08	3.14	*									
NT2RP2001436	3.76	2.25	2.26	8.78	8.61	8.75	5.22	4.80	6.42	**									
NT2RP2001448	3.29	2.41	1.73	3.63	4.88	4.33	2.34	3.35	3.86	*									
NT2RP2001445	2.95	1.26	2.68	2.98	3.78	3.07	2.47	3.15	2.23										
NT2RP2001449	2.88	2.13	1.40	3.15	3.39	4.62	2.6	3.60	1.97										
NT2RP2001458	4.05	2.94	3.13	3.77	4.91	3.85	3.71	4.15	3.13										
NT2RP2001467	2.37	1.91	2.75	5.44	4.55	6.16	5.15	4.88	3.4	**									
NT2RP2001469	10.04	7.34	9.26	5.41	8.75	6.36	6.52	6.42	6.37	*									
NT2RP2001488	6.23	4.15	2.86	6.30	5.94	4.07	6.36	5.86	4.85										
NT2RP2001495	14.26	10.91	10.35	11.90	13.38	11.11	12.39	11.10	12.13										
NT2RP2001499	4.67	3.29	2.95	6.59	7.16	8.76	5.49	6.33	5.02	**									
NT2RP2001506	4.89	3.71	3.86	7.29	8.04	7.88	5.96	6.72	7.88	**									
NT2RP2001508	6.85	6.36	6.72	17.18	14.22	13.59	7.65	11.84	6.81	**									
NT2RP2001511	11.59	6.20	8.17	12.86	12.37	12.22	11.15	9.45	10.6										
NT2RP2001514	6.61	4.54	5.10	6.50	5.89	6.49	5.87	6.17	7.22										
NT2RP2001528	2.37	1.99	2.43	3.12	2.75	2.57	4.35	3.09	2.57										
NT2RP2001526	12.96	5.00	5.99	26.60	29.55	19.41	14.77	8.09	13.41	*									
NT2RP2001529	8.76	6.16	5.20	6.03	7.62	5.55	4.77	5.25	10.17										
NT2RP2001536	3.16	2.19	1.50	3.33	3.18	2.23	2.35	2.79	2.24										
NT2RP2001538	75.84	48.30	57.88	103.08	97.23	96.04	48.73	42.09	45.9	*									
NT2RP2001547	5.37	2.86	3.64	4.76	4.73	4.52	3.5	4.96	4.64										

83/180

1643

1644

NT2RF2001560	6.39	4.64	4.29	5.82	7.13	5.81	3.38	4.66	5.13										
NT2RF2001562	4.89	3.58	3.48	6.44	6.82	4.81	4.71	5.39	5.07										
NT2RF2001566	7.48	4.52	5.51	7.16	5.92	8.75	7.73	7.60	6.5										
NT2RF2001569	14.82	5.79	9.60	21.83	22.56	14.28	10.25	9.70	10.1										
NT2RF2001576	10.55	5.49	5.69	8.15	9.33	7.45	8.98	9.68	8.51										
NT2RF2001581	56.76	28.34	28.83	65.72	65.95	57.58	33.46	29.31	29.57										
NT2RF2001597	6.52	3.84	3.20	6.75	8.45	4.27	5.43	7.30	6.46										
NT2RF2001601	1.39	1.22	0.85	2.84	5.69	3.38	1.83	3.28	2.5	*	+	*	+						
NT2RF2001613	0.98	1.39	1.71	1.95	1.58	2.25	1.57	2.65	2.69										
NT2RF2001628	3.83	3.04	3.39	4.74	7.75	4.57	4.66	5.20	3.94										
NT2RF2001634	9.71	7.65	8.42	9.38	5.92	8.18	7.57	6.78	7.74										
NT2RF2001635	6.36	3.48	2.24	6.23	7.58	4.38	4.88	1.74	2.85										
NT2RF2001648	2.86	2.10	1.03	7.27	5.03	4.32	4.44	3.32	7.02	*	+								
NT2RF2001662	9.75	5.05	6.57	13.09	11.75	8.88	7.01	6.63	7.99										
NT2RF2001663	3.29	2.74	2.56	3.86	4.83	6.87	3.87	4.11	4.21			**	+						
NT2RF2001672	3.92	2.66	2.42	6.76	8.23	7.05	3.9	5.21	5.15	**	+	*	+						
NT2RF2001675	2.35	2.00	2.38	1.25	1.56	1.93	1.59	2.56	2.41	*	-								
NT2RF2001677	6.62	5.40	3.75	5.38	8.63	6.75	8.06	7.03	7.46										
NT2RF2001678	3.81	2.77	2.79	5.76	5.75	5.77	3.78	5.60	5.43	**	+								
NT2RF2001683	1.31	1.34	1.35	2.92	5.85	2.75	1.53	1.74	1.61			**	+						
NT2RF2001699	10.48	4.46	4.39	9.39	8.26	5.63	7.71	4.72	6.45										
NT2RF2001707	6.36	2.69	3.12	4.80	5.89	4.38	5.21	3.89	4.02										
NT2RF2001728	4.31	2.23	2.64	5.76	5.81	5.36	2.53	3.30	4.19	*	+								
NT2RF2001721	5.95	3.63	4.33	4.87	4.91	5.43	4.03	4.62	4.71										
NT2RF2001748	9.64	7.71	6.71	10.42	9.86	6.60	4.64	5.42	6.18										
NT2RF2001748	8.04	6.16	5.85	6.53	8.57	9.79	7.32	7.38	8.28										
NT2RF2001755	8.56	5.19	5.01	5.45	6.63	4.59	3	4.11	4.45										
NT2RF2001762	3.51	1.45	1.56	4.01	2.49	1.10	1.33	1.59	1.38										
NT2RF2001768	10.52	5.70	5.26	8.83	8.48	7.75	7.16	7.38	7.69										
NT2RF2001769	10.19	4.14	4.34	4.02	3.67	3.86	2.04	3.80	3.12										
NT2RF2001784	3.41	2.66	3.05	4.40	6.83	4.24	3.51	4.60	5.21										
NT2RF2001805	8.47	4.44	5.36	7.33	9.55	7.18	6.45	7.26	6.85										
NT2RF2001813	0.85	0.76	1.30	1.56	0.97	1.22	1.03	2.43	0.53										
NT2RF2001817	3.31	2.32	3.38	2.20	3.73	2.38	1.83	3.68	1.91										
NT2RF2001818	9.15	4.97	5.99	7.22	8.04	4.90	5.14	6.97	4.17										
NT2RF2001837	6.67	3.70	3.89	10.21	8.70	8.64	6.67	5.27	5.41	*	+								
NT2RF2001839	8.94	4.07	4.05	8.65	8.01	5.90	7.01	4.33	4.71										
NT2RF2001861	3.92	3.91	2.96	5.38	4.82	4.41	3.85	3.89	4.28	*	+								
NT2RF2001869	3.96	3.68	2.84	5.29	6.76	6.36	4.79	4.96	8.38	**	+								
NT2RF2001876	5.26	4.39	3.67	5.40	6.52	6.44	4.25	3.45	3.89	*	+								
NT2RF2001878	2.96	2.08	2.84	3.77	3.75	3.70	4.02	3.19	4.69	*	+								
NT2RF2001881	3.61	3.23	3.04	4.01	3.35	3.50	1.51	1.79	2.14			**	-						
NT2RF2001883	14.84	8.25	6.92	8.52	8.12	7.84	10.33	7.28	8.44										
NT2RF2001884	13.60	7.36	6.43	4.80	5.47	5.55	7.44	5.61	6.14										
NT2RF2001885	4.58	2.98	2.92	4.56	5.26	4.27	4.8	4.09	3.45										
NT2RF2001898	5.25	3.59	4.61	5.09	5.82	4.63	4.24	6.45	7.13										
NT2RF2001900	3.76	2.05	3.66	6.01	5.52	2.71	3.58	3.82	6.81										
NT2RF2001903	26.77	19.19	22.63	20.41	23.55	21.60	18.49	17.64	17.95										
NT2RF2001907	6.26	4.16	3.66	9.32	10.90	7.90	6.73	6.46	7.99	*	+								
NT2RF2001915	2.75	1.61	1.89	3.01	6.15	2.73	2.2	4.12	4.37										
NT2RF2001921	13.96	7.17	5.50	7.19	5.36	4.44	6.09	4.12	4.96										
NT2RF2001926	2.31	1.57	1.52	6.10	5.30	3.82	6.1	3.59	5.57	*	+	*	+						
NT2RF2001933	7.86	5.07	6.52	8.86	5.68	6.54	7.83	5.74	7.63										
NT2RF2001936	1.63	0.95	0.99	1.17	2.42	2.36	1.86	2.55	1.83										
NT2RF2001940	51.19	30.10	31.53	33.01	35.70	30.61	29.07	28.57	30.35										
NT2RF2001946	3.36	2.65	3.35	3.35	3.83	4.97	4.68	3.30	3.45										

84/180

【0725】

【表254】

1645

1646

NT2RP2001947	4.91	3.61	5.81	3.96	7.73	5.13	4.97	5.37	4.61										
NT2RP2001948	3.08	1.21	4.06	4.99	4.92	1.65	1.37	3.34	8.7										
NT2RP2001949	15.21	7.64	6.12	7.09	9.06	8.60	13.91	9.28	14.64										
NT2RP2001950	8.23	4.55	5.29	5.46	6.80	5.70	8.22	5.90	10.07										
NT2RP2001976	2.14	2.20	2.33	1.64	3.47	2.44	1.48	2.24	2.16										
NT2RP2001978	4.60	3.86	2.35	6.96	6.45	5.14	6.22	4.96	6.39	*	+	*	+						
NT2RP2001985	3.92	3.42	3.57	5.93	6.65	5.91	5.3	5.09	5.9	**	+	**	+						
NT2RP2001991	1.73	1.46	2.57	3.16	4.44	3.93	3.02	3.02	2.07	*	+	*	+						
NT2RP2001997	3.98	3.95	3.94	5.87	6.12	4.91	4.68	4.05	3.66	*	+	*	+						
NT2RP2002015	78.11	51.57	65.21	141.26	146.10	108.68	76.93	62.92	81.97	**	+	**	+						
NT2RP2002017	3.82	3.00	1.73	4.92	6.18	4.74	4	3.36	3.11	*	+	*	+						
NT2RP2002025	9.38	5.00	3.82	6.47	6.74	7.41	7.27	7.03	6.73										
NT2RP2002030	14.24	9.95	8.14	32.58	35.24	33.11	14.46	16.78	20.02	**	+	**	+						
NT2RP2002032	7.60	6.08	6.71	7.52	10.42	7.21	9.78	7.83	10										
NT2RP2002033	10.00	6.88	8.54	14.32	18.25	17.32	8.01	10.19	9.71	**	+	**	+						
NT2RP2002041	1.30	1.42	1.01	2.33	2.65	2.99	2.24	3.22	3.54	**	+	**	+						
NT2RP2002046	2.29	2.31	3.63	4.90	5.83	4.05	4.05	4.50	4.31	*	+	*	+						
NT2RP2002047	5.55	4.39	6.12	3.39	3.21	2.86	3.07	2.96	1.09	*	+	*	+						
NT2RP2002059	8.38	3.98	6.12	10.46	10.43	10.14	8.27	8.23	7.23	*	+	*	+						
NT2RP2002062	6.47	4.41	3.60	6.50	9.32	5.86	4.66	4.62	6.58										
NT2RP2002058	3.62	2.82	3.02	3.46	3.52	2.23	2.78	3.89	2.56										
NT2RP2002060	6.58	3.14	4.55	4.58	5.81	5.66	5.55	7.36	5.35										
NT2RP2002063	1.56	1.90	1.51	3.69	1.67	1.86	2.22	2.63	1.71										
NT2RP2002066	5.03	3.37	4.61	4.73	5.21	5.32	7.33	6.17	4.62										
NT2RP2002078	0.79	0.79	0.34	1.28	2.20	1.05	0.97	2.47	0.94										
NT2RP2002076	3.86	2.57	2.52	3.36	3.56	2.78	2.73	4.09	2.15										
NT2RP2002078	5.54	3.35	3.42	13.66	10.39	8.08	7.93	6.64	6.4	*	+	*	+						
NT2RP2002079	5.14	3.23	1.70	5.80	4.94	6.51	3.67	4.05	3.99										
NT2RP2002099	7.45	3.48	2.47	4.21	4.13	3.43	3.32	4.93	4.92										
NT2RP2002105	5.64	3.25	3.05	3.88	4.16	3.68	4.68	5.62	4.37										
NT2RP2002115	0.92	0.69	0.55	1.83	1.20	1.32	0.97	2.15	0.81	*	+	*	+						
NT2RP2002124	2.28	1.30	1.91	4.70	4.64	3.30	3.98	3.75	2.5	*	+	*	+						
NT2RP2002137	2.93	1.88	1.87	2.18	3.16	2.61	3.4	4.11	2.95	*	+	*	+						
NT2RP2002139	4.33	3.54	3.42	3.56	4.04	4.02	5.23	4.66	5.13	*	+	*	+						
NT2RP2002154	5.53	2.76	1.92	4.83	6.57	3.88	4.83	4.72	5.4										
NT2RP2002155	279.79	155.93	163.22	227.28	242.49	184.60	219.6	179.59	177.9										
NT2RP2002172	4.14	2.59	2.22	3.81	3.52	4.02	3.34	4.90	3.32										
NT2RP2002185	4.32	3.52	2.95	4.55	4.64	4.41	4.65	5.42	5.45										
NT2RP2002188	11.41	5.54	8.75	9.54	13.32	9.41	7.96	10.55	9.63										
NT2RP2002192	3.64	3.48	3.53	4.30	3.68	3.71	1.91	3.83	2.29										
NT2RP2002193	3.15	2.72	2.77	3.68	4.01	3.41	3.89	3.36	4.16	*	+	*	+						
NT2RP2002206	2.07	2.36	2.72	6.19	4.41	5.19	4.33	5.08	2.51	**	+	**	+						
NT2RP2002219	4.17	1.29	1.62	2.78	4.30	2.60	1.31	1.97	1.84										
NT2RP2002231	2.75	2.39	1.20	3.02	3.57	1.95	2.15	1.47	2.21										
NT2RP2002232	5.59	1.67	2.23	5.04	5.05	3.28	3.82	4.55	3.16										
NT2RP2002234	7.15	4.93	3.90	3.84	3.33	2.89	4.86	6.74	5.47										
NT2RP2002239	23.74	15.37	16.41	23.91	26.96	19.68	8.99	12.98	10.06										
NT2RP2002252	9.96	4.94	5.61	5.48	5.08	6.39	5.19	6.06	5.88										
NT2RP2002256	1.33	1.22	1.37	1.71	2.37	2.14	1.73	2.95	1.47	*	+	*	+						
NT2RP2002257	2.29	1.76	1.74	4.11	5.09	2.83	4.04	4.42	3.81	*	+	*	+						
NT2RP2002259	3.72	2.30	2.90	6.32	3.45	2.90	3.06	3.48	1.66										
NT2RP2002264	2.47	1.33	1.14	6.07	7.37	5.74	2.09	3.51	3.03	**	+	**	+						
NT2RP2002267	8.31	4.57	4.68	12.99	14.87	10.14	12.21	9.31	10.07	*	+	*	+						
NT2RP2002278	7.39	4.62	5.64	7.88	7.73	8.65	3.38	3.42	4.07										
NT2RP2002281	8.20	4.58	6.60	7.60	8.32	8.02	5.18	6.11	4.33										
NT2RP2002288	5.39	5.46	4.44	3.41	3.45	3.50	3.57	3.54	3.96	**	+	**	+						

85/180

【0726】

【表255】

1647

1648

NT1RF2002391	13.36	8.93	10.00	7.24	12.33	7.03	8.51	6.90	8.43				
NT1RF2002399	4.86	3.21	3.87	7.31	5.99	7.44	5.79	6.94	6.46	*	+	*	+
NT1RF2002394	3.12	1.09	1.07	3.72	6.64	4.48	2.39	2.10	2.14	*	+		
NT1RF2002312	3.00	2.02	1.91	4.87	5.25	3.26	3.11	3.70	3.89	*	+	*	+
NT1RF2002316	2.57	2.29	2.38	6.74	6.43	5.78	3.23	3.23	4.39	**	+	*	+
NT1RF2002325	2.17	2.03	1.50	3.32	3.39	2.92	1.65	3.11	3.18	**	+		
NT1RF2002333	6.45	4.83	4.75	7.88	10.32	7.81	5.66	5.80	6.3	*	+		
NT1RF2002371	4.90	4.23	3.63	9.29	8.56	8.23	9.75	10.58	7.26	**	+	**	+
NT1RF2002373	5.37	4.02	2.70	5.83	10.05	6.25	5.7	8.27	6.72				
NT1RF2002381	0.73	0.29	0.85	0.79	0.90	2.57	1.16	2.65	1.41				
NT1RF2002385	7.34	2.40	2.24	6.24	3.86	3.39	5.09	3.89	4.74				
NT1RF2002394	1.71	0.33	0.18	1.03	1.49	1.31	0.28	1.27	2.19				
NT1RF2002408	2.38	1.66	1.45	4.45	2.73	2.67	1.95	4.44	3.16				
NT1RF2002409	29.85	16.62	15.12	29.12	39.51	28.40	19.16	20.28	16.99				
NT1RF2002424	3.78	2.45	1.98	3.14	4.67	3.25	3.81	5.82	3.46				
NT1RF2002426	5.16	3.36	3.05	8.68	9.29	8.07	5.5	8.86	7.03	**	+	*	+
NT1RF2002429	6.36	5.02	5.09	9.72	12.33	8.37	9.84	17.67	16.81	*	+	*	+
NT1RF2002437	3.49	2.56	3.29	4.17	7.17	4.10	3.26	6.17	5.32				
NT1RF2002439	11.07	5.27	5.30	11.81	8.46	7.22	11.52	9.36	7.78				
NT1RF2002443	6.40	2.74	3.03	4.62	5.05	4.46	4.75	2.98	3.74				
NT1RF2002457	2.28	2.49	1.70	3.54	4.01	3.48	4.07	3.72	3.08	**	+	*	+
NT1RF2002464	5.19	2.78	3.13	3.90	4.79	4.00	5.08	3.74	4				
NT1RF2002475	3.58	3.74	3.05	8.04	7.22	4.99	7.48	6.02	7.62	*	+	**	+
NT1RF2002479	3.49	2.33	2.32	3.60	4.32	2.72	2.92	2.66	5.14				
NT1RF2002487	4.86	2.73	2.49	4.04	4.25	4.00	3.16	3.11	3.07				
NT1RF2002498	2.48	0.99	1.21	3.47	2.96	2.53	1.35	1.52	1.58				
NT1RF2002503	13.02	6.05	8.78	12.14	16.89	12.87	9.04	8.81	7.66				
NT1RF2002504	6.63	3.00	4.84	4.05	6.27	4.67	6.68	4.71	5.18				
NT1RF2002510	15.40	9.87	11.00	12.38	17.28	17.15	18.56	12.92	13.19				
NT1RF2002528	1.61	1.78	1.33	4.08	3.77	4.83	3.97	4.73	4.31	**	+	**	+
NT1RF2002527	11.26	7.87	9.14	12.36	15.57	11.93	8.08	6.87	9.06				
NT1RF2002533	15.80	10.32	13.55	16.21	16.47	14.65	18.71	12.94	18.73				
NT1RF2002537	6.78	4.47	5.46	7.12	8.21	8.66	4.34	3.85	6.54	*	+		
NT1RF2002542	11.84	6.86	7.87	24.97	24.70	21.27	12.25	9.81	10.65	**	+		
NT1RF2002546	3.51	1.75	1.39	2.49	2.71	2.52	4.4	3.54	3.7				
NT1RF2002549	8.05	4.99	5.19	5.57	6.51	7.45	6.2	3.49	5.35				
NT1RF2002564	13.08	7.54	8.36	11.61	12.09	10.41	11.1	8.10	13.89				
NT1RF2002591	9.73	4.99	4.71	11.69	11.90	10.05	7.9	7.42	7.09				
NT1RF2002595	5.43	4.01	5.43	9.33	7.85	7.01	6.61	6.19	7.33	*	+	*	+
NT1RF2002603	4.82	4.74	4.84	5.43	11.27	8.16	5.69	6.45	7.55			*	+
NT1RF2002606	5.86	3.02	3.06	8.03	9.33	3.93	3.99	4.72	6.99				
NT1RF2002609	4.71	2.92	3.43	5.18	4.82	3.59	3.34	4.09	4.4				
NT1RF2002618	4.82	3.33	2.74	6.13	4.63	4.67	4.95	4.51	4.42				
NT1RF2002621	10.26	6.84	5.48	15.22	14.98	13.05	11.07	8.62	10.72	*	+		
NT1RF2002643	4.22	2.96	3.21	5.73	8.43	4.77	4.53	4.98	4.94			*	+
NT1RF2002672	4.36	3.45	3.37	8.96	12.04	8.60	8.5	8.50	11.85	**	+	**	+
NT1RF2002673	2.97	2.38	1.11	7.44	9.35	7.43	5.4	7.46	8.29	**	+	**	+
NT1RF2002674	1.07	1.16	1.07	0.86	1.66	1.60	1.52	2.10	1.72			*	+
NT1RF2002686	3.43	3.39	4.42	4.11	5.80	4.25	4.81	4.16	5.05				
NT1RF2002698	13.80	10.26	10.99	17.41	16.88	13.34	9.74	11.51	8.03				
NT1RF2002695	6.80	3.06	3.92	5.81	7.30	4.59	7.03	4.43	5.61				
NT1RF2002701	6.95	4.89	4.37	8.51	9.98	9.57	8.2	7.53	9.25	*	+	*	+
NT1RF2002706	4.89	2.72	3.50	5.60	7.16	5.82	4.64	4.20	5.97	*	+		
NT1RF2002719	42.99	27.04	33.49	47.10	50.09	39.86	54.16	44.21	55.65			*	+
NT1RF2002721	7.76	5.23	6.54	9.71	10.64	8.40	10.35	8.03	8.87	*	+		
NT1RF2002727	0.98	1.45	0.99	2.09	1.15	2.50	2.3	2.04	1.73			*	+

86/180

【0727】

【表256】

1649

1650

NT2RF2002734	4.53	3.02	5.80	12.41	12.84	10.69	6.86	8.07	7.85	**	+	*	+
NT2RF2002736	3.63	2.27	2.67	2.07	2.02	2.04	2.87	2.60	2.01				
NT2RF2002740	2.59	1.02	0.94	3.18	2.63	2.29	2.78	2.96	1.96				
NT2RF2002741	5.52	4.27	3.15	7.73	8.99	8.94	4.51	5.06	7.43	**	+		
NT2RF2002759	7.28	6.29	4.77	14.35	17.57	18.80	8.32	9.26	7.61	**	+		
NT2RF2002753	11.68	7.46	7.74	12.78	17.74	15.50	11.31	10.02	12.22	*	+		
NT2RF2002753	11.55	5.48	11.53	10.53	6.13	11.57	7.42	7.93	9.43				
NT2RF2002760	8.78	4.40	4.62	7.89	8.63	6.01	6.34	6.38	7.33				
NT2RF2002769	3.29	2.63	2.68	3.72	6.64	6.67	2.86	4.11	3.55	*	+		
NT2RF2002778	9.07	6.03	9.70	7.44	6.87	7.92	6.93	7.76	4.98				
NT2RF2002791	6.58	4.82	4.00	9.50	14.75	9.25	8.23	6.79	7.02	*	+		
NT2RF2002800	6.57	4.20	5.63	10.46	11.33	12.38	5.4	8.07	7.04	**	+		
NT2RF2002805	1.48	1.18	0.66	2.57	1.66	1.18	2.89	3.53	1.96		*	+	
NT2RF2002811	5.70	5.54	4.77	8.54	7.13	7.69	6.53	7.67	6.08	**	+		
NT2RF2002824	9.12	5.93	7.91	13.68	13.22	9.65	9.82	10.22	11.6	*	+		
NT2RF2002839	3.89	2.03	2.96	3.87	4.52	3.28	3.17	3.43	3.41				
NT2RF2002845	2.29	1.84	1.77	4.04	4.31	4.72	3.6	4.26	3.16	**	+	**	+
NT2RF2002857	0.99	1.45	1.80	1.98	2.27	1.78	2.36	3.14	1.89				
NT2RF2002862	11.21	6.20	5.58	10.84	12.86	10.44	6.99	7.12	10.71				
NT2RF2002880	5.70	4.03	2.74	3.50	4.84	3.87	4.05	5.72	5				
NT2RF2002885	6.90	4.59	4.82	5.83	6.45	4.16	3.34	4.76	3.08				
NT2RF2002891	5.76	3.80	3.33	5.44	6.69	6.13	4.92	4.49	5.35				
NT2RF2002907	4.12	1.98	2.30	4.77	3.91	2.49	2.25	3.24	2.04				
NT2RF2002925	3.23	2.04	2.18	4.98	4.44	5.21	3.38	2.81	4.67	**	+		
NT2RF2002927	14.45	8.55	11.84	14.25	14.86	13.10	10.66	9.50	13.04				
NT2RF2002928	1.42	1.26	2.32	3.26	2.52	3.14	1.44	1.91	1.88	*	+		
NT2RF2002929	6.54	3.13	3.18	6.60	7.00	5.63	5.25	5.85	5.87				
NT2RF2002934	5.87	2.70	3.00	3.46	2.95	4.09	3.58	3.88	3.47				
NT2RF2002939	6.87	3.02	3.14	4.78	4.45	4.28	3.95	4.36	3.63				
NT2RF2002942	4.16	2.79	3.25	6.95	8.21	6.01	4.14	5.76	4.58	**	+		
NT2RF2002954	3.73	2.07	3.02	3.75	4.03	3.04	2.28	3.89	5.22				
NT2RF2002959	5.43	4.36	4.62	6.19	7.91	6.08	3.63	5.75	5.03	*	+		
NT2RF2002974	2.77	2.53	1.82	5.32	4.88	3.20	3.66	3.70	3.24	*	+	*	+
NT2RF2002976	1.81	1.66	2.46	4.07	3.02	2.77	2.16	2.65	2.13	*	+		
NT2RF2002979	10.96	6.09	6.26	13.05	14.90	10.76	8.18	9.68	7.32				
NT2RF2002980	8.71	5.49	6.33	14.65	15.05	11.66	8.24	9.16	9.26	**	+		
NT2RF2002986	8.28	6.07	5.22	8.21	6.48	6.46	9.09	7.74	9.39				
NT2RF2002987	6.13	3.28	3.28	8.77	8.51	7.89	4.85	7.00	9.15	*	+		
NT2RF2002988	34.52	23.01	24.20	21.24	19.88	21.98	15.82	15.65	16.56		*	-	
NT2RF2002993	4.35	3.19	4.08	2.57	3.44	2.83	3.21	3.84	2.8				
NT2RF2003000	6.81	5.24	5.01	12.83	14.50	14.13	6.77	6.65	8.42	**	+		
NT2RF2003008	3.03	1.86	2.21	2.77	3.21	3.26	2.46	3.49	5.58				
NT2RF2003020	7.91	3.15	3.03	14.51	13.63	11.55	10.67	9.71	9.8	**	+	*	+
NT2RF2003032	4.25	3.36	3.04	5.65	7.30	4.26	5.14	2.86	5.02				
NT2RF2003034	8.64	4.19	5.82	12.73	13.68	11.86	9.6	7.30	8.21	**	+		
NT2RF2003042	3.77	2.17	2.53	3.68	4.54	3.65	3.09	3.66	3.89				
NT2RF2003050	2.09	1.93	2.12	2.58	4.04	3.16	2.04	3.12	2.84	*	+		
NT2RF2003060	6.89	6.04	6.20	6.11	6.61	6.02	4.64	5.08	4.58		**	-	
NT2RF2003073	5.10	4.79	4.81	10.73	11.79	9.58	6.83	8.25	4.87	**	+		
NT2RF2003099	3.77	3.26	2.86	5.64	6.07	7.01	4.27	4.57	5.19	**	+	*	+
NT2RF2003108	3.73	1.70	0.71	4.43	4.78	3.41	2.53	3.13	3.98				
NT2RF2003115	12.63	7.03	6.49	10.94	5.30	4.75	6.51	5.99	10.91				
NT2RF2003117	9.96	4.65	5.66	15.04	15.48	11.38	8.83	7.84	6.17	*	+		
NT2RF2003121	3.53	2.40	1.92	4.90	5.00	3.18	3.72	4.52	4.42		*	+	
NT2RF2003125	5.32	2.20	2.34	3.41	4.18	3.51	3.6	4.31	3.11				
NT2RF2003127	3.09	3.27	3.35	3.25	3.63	2.74	2.3	4.46	3.68				

87/180

1651

1652

NTZRF2003129	3.68	2.64	1.93	5.72	5.89	5.75	3.03	4.40	2.82	**	+		
NTZRF2003137	2.40	2.79	2.71	6.74	6.38	5.76	4.22	6.41	4.31	**	+	*	+
NTZRF2003138	6.42	2.67	2.97	5.99	6.92	3.98	5.12	3.06	1.92				
NTZRF2003146	4.44	2.51	1.78	3.73	3.26	2.77	3.76	2.57	1.66				
NTZRF2003148	9.10	6.45	5.51	11.73	13.86	11.19	8.71	8.13	7.46	*	+		
NTZRF2003150	3.26	2.20	1.35	8.65	2.99	4.86	3.92	2.84	8.35				
NTZRF2003157	7.49	3.86	3.67	8.41	10.43	9.55	4.96	6.45	5.87	*	+		
NTZRF2003158	1.88	1.89	2.17	2.26	3.00	2.46	2.43	2.76	2.85			*	+
NTZRF2003161	1.04	1.33	0.76	2.12	4.38	4.18	1.59	2.84	8.91	*	+		
NTZRF2003164	2.83	1.78	1.70	2.90	2.78	2.57	2.53	2.97	2.44				
NTZRF2003165	4.31	2.10	2.06	5.98	4.84	6.84	5.12	3.81	4.72	*	+		
NTZRF2003177	3.18	2.52	2.22	3.53	2.99	3.63	4.35	2.80	2.79				
NTZRF2003179	4.54	3.39	3.36	5.90	7.70	7.29	4.85	4.79	6.24	**	+		
NTZRF2003194	16.94	9.59	9.74	7.86	8.77	6.84	7.23	6.50	9.93				
NTZRF2003206	0.19	0.73	0.54	2.02	2.10	1.11	1.07	1.15	1.17	*	+	*	+
NTZRF2003210	5.52	2.50	2.65	2.94	4.61	3.60	3.44	3.99	4.15				
NTZRF2003227	2.55	1.52	2.78	3.96	4.66	3.48	2.52	3.60	4.44	*	+		
NTZRF2003228	5.50	4.11	4.96	4.07	4.64	3.51	3.63	3.86	2.66				
NTZRF2003230	1.04	1.41	1.38	3.75	3.72	3.44	8.77	4.96	7.21	**	+	**	+
NTZRF2003231	6.83	5.52	4.87	9.61	7.64	6.47	5.75	5.89	8.09				
NTZRF2003237	4.46	2.56	2.35	5.51	7.13	6.33	3.56	4.31	3.67	*	+		
NTZRF2003239	4.50	2.01	3.71	6.44	6.32	5.76	4.01	4.23	4.42	*	+		
NTZRF2003243	5.46	3.20	3.57	7.44	6.11	7.58	5.91	6.40	3.87	*	+		
NTZRF2003265	5.61	3.24	3.60	7.47	8.92	7.01	5.38	4.10	6.74	*	+		
NTZRF2003267	3.97	3.06	3.71	7.15	8.86	6.88	4.28	4.40	5.84	**	+		
NTZRF2003272	5.37	3.98	5.63	6.49	6.56	6.62	7.54	6.51	7.61	*	+	*	+
NTZRF2003277	9.14	5.91	4.66	7.52	10.35	9.11	9.97	7.77	15.8				
NTZRF2003280	3.01	2.25	1.41	4.02	6.71	7.68	6.13	4.20	7.59	*	+	*	+
NTZRF2003286	3.53	1.84	2.37	2.62	3.15	2.83	2.96	2.70	4.01				
NTZRF2003293	6.85	4.64	6.03	12.22	12.54	11.97	6.66	5.15	8.8	**	+		
NTZRF2003295	4.81	3.25	3.18	3.96	8.36	5.27	4.16	4.98	3				
NTZRF2003297	1.97	1.06	1.42	2.82	3.09	2.49	1.97	1.89	1.68	*	+		
NTZRF2003300	5.99	4.89	4.68	7.75	7.40	7.47	7.28	9.19	9.08	**	+	*	+
NTZRF2003302	4.65	3.24	4.39	8.90	10.20	7.29	4.36	7.27	5.11	**	+		
NTZRF2003307	1.67	1.09	0.57	2.24	1.67	2.40	2.82	1.84	1.76				
NTZRF2003308	3.09	2.17	1.85	4.09	5.19	2.83	3.04	2.74	3.16				
NTZRF2003311	6.85	3.58	2.13	4.65	6.66	4.36	3.88	3.65	4.23				
NTZRF2003329	3.07	1.86	1.87	3.19	5.07	3.49	3.77	3.82	5.96				
NTZRF2003339	2.38	1.55	1.29	2.90	3.98	3.91	2.69	3.47	2.24	*	+		
NTZRF2003345	1.83	1.44	1.40	1.51	1.52	1.92	2.28	2.65	1.28				
NTZRF2003347	1.48	2.10	1.67	2.03	5.75	1.76	2.44	3.10	4.09		*	+	
NTZRF2003367	1.26	0.98	1.42	1.39	1.59	1.55	1.21	2.14	1.04				
NTZRF2003369	3.82	2.31	1.37	1.42	2.10	1.87	3.19	2.85	1.99				
NTZRF2003383	7.18	3.57	4.41	16.30	14.96	15.98	8.79	9.62	11.29	**	+	*	+
NTZRF2003390	9.92	6.14	6.73	11.71	12.19	9.52	7.92	9.43	8.34				
NTZRF2003391	35.23	21.64	23.59	36.95	36.23	27.51	23.69	17.29	17.85				
NTZRF2003393	2.40	1.57	1.83	4.13	5.18	3.56	3.96	4.34	3.87	*	+	**	+
NTZRF2003394	4.02	2.41	2.76	12.16	9.99	10.68	6.12	6.15	3.96	**	+		
NTZRF2003401	2.33	1.80	1.86	3.02	4.68	2.41	3.02	4.51	3.57		*	+	
NTZRF2003403	1.23	1.40	1.41	3.20	3.23	4.51	3.04	3.80	3.41	**	+	**	+
NTZRF2003433	8.96	4.52	3.52	6.71	5.66	5.39	7.4	6.01	5.01				
NTZRF2003446	3.20	3.09	2.41	6.94	6.16	6.94	13.01	11.43	14.04	**	+	**	+
NTZRF2003446	5.05	4.02	2.72	4.09	6.31	3.82	5.45	4.95	5.35				
NTZRF2003456	4.21	2.96	2.69	10.80	8.14	8.43	6.15	5.44	4.71	**	+	*	+
NTZRF2003466	5.26	3.68	3.82	5.95	5.44	4.60	3.82	5.23	9				
NTZRF2003469	3.53	2.12	2.45	3.89	4.69	5.28	2.75	4.01	3.09	*	+		

88/180

1653

1654

NT2RP2003478	11.59	7.42	9.22	28.44	23.50	24.05	11.29	12.07	8.19	**	+		
NT2RP2003471	0.62	0.28	0.53	1.86	1.08	1.71	2.23	2.31	0.86	*	+		
NT2RP2003480	15.63	7.31	7.47	13.91	14.92	13.14	9.58	7.59	11				
NT2RP2003495	8.78	5.33	4.65	3.96	5.20	6.08	4.27	5.58	4.14				
NT2RP2003499	3.16	1.30	1.31	2.42	1.62	2.16	3.79	4.26	2.53				
NT2RP2003505	2.95	2.52	1.64	4.06	3.25	3.65	2.65	3.70	2.81	*	+		
NT2RP2003506	4.36	2.44	2.89	4.61	6.57	3.32	3.86	4.37	5.74				
NT2RP2003511	5.80	4.98	5.36	9.63	8.04	5.73	6.43	6.77	8.36			*	+
NT2RP2003513	3.23	2.52	3.10	3.94	3.00	3.76	2.27	3.48	3.18				
NT2RP2003517	1.52	0.95	2.01	2.87	2.13	1.37	2.66	3.16	3.17			*	+
NT2RP2003522	21.16	8.31	12.55	21.51	17.78	15.40	9.2	5.69	8.01				
NT2RP2003525	6.58	6.05	5.00	12.44	12.64	12.83	8.86	7.54	7.95	**	+	*	+
NT2RP2003533	7.73	4.59	4.51	11.94	12.52	10.34	6.62	8.25	8.72	**	+		
NT2RP2003541	9.89	7.73	6.72	8.34	7.49	6.40	6.78	6.83	5.85				
NT2RP2003543	4.46	3.26	2.49	5.01	7.76	4.19	6.57	7.85	7.39			**	+
NT2RP2003545	6.37	3.24	4.48	2.58	2.60	1.65	1.96	3.63	2.3				
NT2RP2003549	1.78	1.16	2.25	3.59	3.08	3.14	2.24	2.88	3.16	*	+		
NT2RP2003564	1.63	1.70	1.81	2.44	3.74	2.88	2.97	3.23	1.66	*	+		
NT2RP2003565	9.14	3.08	4.12	8.63	10.17	6.24	4.03	4.24	3.56				
NT2RP2003567	7.44	5.21	4.96	7.20	9.00	7.04	7.75	6.53	4.86				
NT2RP2003575	5.24	1.86	2.00	2.78	2.67	1.70	1.73	2.24	4.67				
NT2RP2003576	208.36	132.21	112.56	100.63	118.10	86.36	71.48	50.82	50.69			*	-
NT2RP2003579	56.28	38.17	48.67	28.49	15.58	24.16	19.34	17.93	21.34	*	-	**	-
NT2RP2003581	4.71	3.22	3.45	3.09	5.04	4.47	3.46	3.82	4.77				
NT2RP2003587	8.55	4.99	7.99	8.79	9.50	8.44	7.38	8.78	13.4				
NT2RP2003590	11.27	7.70	8.07	4.15	4.86	4.77	3.73	6.36	4.84	*	-	*	-
NT2RP2003593	9.63	4.82	5.47	13.80	9.73	5.79	6.89	8.08	6.91				
NT2RP2003596	3.20	2.89	2.89	6.00	8.78	7.99	4.62	4.90	7.08	**	+	*	+
NT2RP2003599	8.81	5.81	5.81	8.37	10.49	10.48	10.61	8.00	12.61				
NT2RP2003600	3.15	1.54	2.36	3.63	5.05	4.21	2.91	3.54	3.28	*	+		
NT2RP2003604	8.61	4.63	5.27	5.66	7.11	7.00	5.84	5.70	5.33				
NT2RP2003629	0.93	0.41	0.97	1.80	1.56	1.57	0.76	2.29	1.4	*	+		
NT2RP2003630	3.31	2.56	2.95	6.23	8.50	6.34	5.52	5.72	4.54	**	+	**	+
NT2RP2003643	16.50	10.48	12.66	12.59	15.91	12.75	9.42	11.38	10.03				
NT2RP2003655	4.54	2.17	1.95	4.91	4.47	3.19	4.31	4.99	6.38				
NT2RP2003664	7.29	4.58	3.44	9.78	13.11	10.33	7.53	12.65	18.19	*	+		
NT2RP2003668	7.64	3.93	2.99	7.77	11.11	7.27	3.61	4.49	4.92				
NT2RP2003687	3.50	2.00	2.53	2.44	3.28	2.52	1.34	3.20	1.86				
NT2RP2003691	3.51	2.23	2.36	4.83	5.26	4.14	2.6	3.93	3.34	*	+		
NT2RP2003702	4.72	3.23	2.97	5.75	5.42	5.03	3.29	5.65	2.48	*	+		
NT2RP2003704	3.03	1.02	1.33	3.00	4.19	2.96	1.48	4.19	2.8				
NT2RP2003706	0.54	0.54	0.40	1.92	1.23	0.53	1.37	2.50	2.1			*	+
NT2RP2003713	3.77	2.04	1.68	4.89	3.40	3.69	3.54	1.79	2.29				
NT2RP2003714	16.93	11.05	8.85	15.34	13.25	10.73	6.94	5.43	4.92				
NT2RP2003727	9.17	5.59	4.98	8.92	8.98	7.11	5.82	4.15	6.45				
NT2RP2003737	4.49	2.62	2.06	3.80	4.50	3.26	2.92	3.29	5.35				
NT2RP2003751	0.82	0.97	1.07	1.33	1.62	0.98	1.33	0.88	0.72				
NT2RP2003760	3.61	2.60	1.42	4.28	5.22	4.19	4.75	3.97	7.45	*	+		
NT2RP2003764	4.43	3.65	3.32	3.81	3.64	3.20	3.86	3.12	8.26				
NT2RP2003769	3.03	1.62	1.45	3.28	5.14	3.51	3.96	2.62	2.26				
NT2RP2003770	11.88	6.14	3.72	10.96	9.10	9.34	9.86	5.90	7.19				
NT2RP2003777	8.28	5.95	4.45	12.14	8.07	7.31	6.16	4.05	5.91				
NT2RP2003781	6.93	4.17	4.88	6.60	9.83	10.25	6.27	5.64	6.39				
NT2RP2003785	5.07	3.24	3.30	5.65	5.69	5.57	6.33	7.73	14.42	*	+		
NT2RP2003793	9.26	6.02	4.92	6.26	7.16	5.41	4.28	4.76	5.16				
NT2RP2003806	6.44	4.78	6.02	12.68	12.04	12.13	5.52	7.88	5.99	**	+		

1655

1656

NT2RF2003825	9.16	5.63	6.57	17.27	18.54	12.04	6.67	8.08	14.03	*	+		
NT2RF2003848	10.64	4.89	5.66	8.31	7.78	5.93	7.12	5.91	8.06				
NT2RF2003857	12.72	6.86	6.25	8.31	8.84	9.18	7.95	6.05	8.74				
NT2RF2003859	6.93	3.73	2.73	12.12	10.40	13.45	3.71	3.90	6.36	**	+		
NT2RF2003871	3.42	3.01	2.13	9.67	10.18	8.65	5.24	4.53	5.97	**	+	*	+
NT2RF2003876	7.74	4.51	4.43	5.67	8.07	7.43	4.37	5.53	5.6				
NT2RF2003878	4.47	2.22	2.10	3.89	4.71	3.64	3.95	3.56	4.06				
NT2RF2003883	5.69	2.59	2.76	3.73	7.92	5.39	4.25	4.87	6.01				
NT2RF2003899	10.09	7.67	7.33	11.75	12.18	9.73	5.01	8.03	5.65				
NT2RF2003902	10.41	8.37	6.78	8.14	9.71	9.88	7.68	5.42	8.06				
NT2RF2003913	13.81	9.98	7.42	16.63	17.90	13.52	10.9	14.66	13.18				
NT2RF2003931	3.74	1.68	1.44	2.28	2.88	2.54	2.24	1.94	2.65				
NT2RF2003948	18.24	10.75	11.51	44.72	39.79	24.81	16	14.58	19.02	*	+		
NT2RF2003959	3.98	2.45	3.31	3.52	4.06	3.60	3	3.05	3.52				
NT2RF2003952	5.00	3.18	4.24	4.00	4.74	3.20	2.55	2.62	4.33				
NT2RF2003968	13.52	6.81	6.24	9.83	14.58	9.98	4.25	4.49	10.21				
NT2RF2003976	5.76	3.40	2.77	10.86	15.30	22.19	5.6	7.99	7.6	*	+		
NT2RF2003981	5.81	3.89	2.20	4.65	4.94	4.43	4.88	3.67	4.28				
NT2RF2003984	11.22	7.15	6.30	8.47	13.43	9.96	9.18	9.47	16.24				
NT2RF2003986	11.50	5.47	4.61	14.29	15.56	15.24	7.95	7.99	8.32	*	+		
NT2RF2003988	5.84	4.44	3.08	11.21	13.07	8.96	7.35	4.72	6.91	**	+		
NT2RF2004013	19.46	11.40	12.00	20.33	26.92	19.32	8.59	11.59	12.13				
NT2RF2004014	5.88	5.77	8.06	11.00	14.73	13.84	6.02	5.49	4.74	**	+		
NT2RF2004036	4.76	2.41	3.64	4.63	4.19	5.70	3.7	3.95	3.26				
NT2RF2004041	2.79	3.61	3.30	4.01	6.06	4.15	3.2	4.29	4.43				
NT2RF2004042	4.23	3.45	2.82	4.59	3.59	5.00	3.97	2.94	3.64				
NT2RF2004049	5.52	3.09	3.20	5.68	4.82	4.18	3.14	3.78	3.4				
NT2RF2004060	6.54	4.19	4.75	5.31	7.44	5.90	6.84	5.31	6.57				
NT2RF2004066	7.64	3.57	3.11	8.07	8.17	6.09	3.54	4.23	4.08				
NT2RF2004069	2.46	2.35	2.84	3.73	4.30	3.52	3.02	4.14	4.07	**	+	*	+
NT2RF2004076	1.40	1.15	1.26	2.49	2.65	1.93	1.27	2.46	1.33	**	+		
NT2RF2004080	2.70	2.23	2.55	3.88	5.93	4.96	4.18	5.58	4.25	*	+	*	+
NT2RF2004081	2.74	2.99	2.36	3.72	4.51	3.72	1.45	3.28	1.61	*	+		
NT2RF2004098	10.83	5.42	4.87	10.62	9.37	7.52	6.04	4.69	6.05				
NT2RF2004108	15.24	8.74	6.82	24.00	21.97	22.21	10.22	12.30	14.43	**	+		
NT2RF2004124	5.29	4.13	3.63	5.87	5.42	5.25	4.18	2.84	4.23				
NT2RF2004130	9.77	7.17	7.05	9.85	13.14	10.78	12.57	13.32	11.04	*	+		
NT2RF2004133	11.24	7.82	7.31	10.46	12.30	8.54	8.71	9.42	8.83				
NT2RF2004141	4.33	2.78	3.55	5.05	6.27	4.10	3.83	4.25	5.14				
NT2RF2004143	3.53	1.25	3.26	3.70	5.10	5.11	2.84	4.94	3.66				
NT2RF2004153	2.68	1.78	2.43	4.24	5.04	5.23	2.05	2.34	1.5	**	+		
NT2RF2004165	21.03	8.19	8.39	7.87	8.05	7.98	5.38	6.52	6.22				
NT2RF2004170	7.13	4.37	2.78	6.23	7.89	6.07	5.24	5.06	3.73				
NT2RF2004172	3.69	2.25	1.50	2.50	3.71	2.71	2.83	3.52	1.97				
NT2RF2004176	7.84	4.13	3.67	5.48	5.12	4.33	5.56	7.38	6.12				
NT2RF2004179	6.87	2.52	2.41	5.35	4.30	3.84	3.98	4.72	4.2				
NT2RF2004187	3.69	2.64	1.86	5.62	6.94	5.86	3.38	4.90	4.03	**	+		
NT2RF2004190	2.07	2.03	2.45	3.29	3.28	2.78	5.06	5.55	4.18	*	+	**	+
NT2RF2004194	6.67	3.78	5.18	7.29	8.60	7.46	5.61	6.42	7				
NT2RF2004196	20.28	5.85	8.53	16.34	14.05	15.75	7.78	7.99	8.4				
NT2RF2004285	10.63	6.42	6.10	11.21	13.23	11.22	6.53	6.15	7.63				
NT2RF2004287	4.42	3.24	2.70	3.44	4.24	3.84	3.13	3.26	3.82				
NT2RF2004326	4.97	4.89	4.35	4.76	5.20	4.65	3.73	3.67	3.35	**	-		
NT2RF2004332	2.49	1.77	2.98	3.76	4.69	3.30	2.85	3.10	2.15	*	+		
NT2RF2004339	4.49	3.56	3.79	6.17	7.37	6.14	4.15	5.46	4.58	**	+		
NT2RF2004340	6.30	3.45	4.77	13.34	11.74	9.18	6.02	6.36	6.66	*	+		

90/180

【0731】

【表260】

1658

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91/180

【表 261】

1659

1660

NT1RF2004709	5.18	3.25	1.93	12.66	12.56	10.94	5.12	4.16	3.85	**	+		
NT1RF2004710	5.83	4.70	2.80	7.69	7.61	6.76	4.34	3.44	4.54	*	+		
NT1RF2004721	11.13	7.44	7.40	6.68	9.65	8.99	11.35	9.52	13.55				
NT1RF2004736	6.91	5.30	5.26	8.14	9.36	7.77	6.53	5.39	5.85	**	+		
NT1RF2004743	2.77	1.82	1.63	5.65	6.03	4.15	4.87	6.71	5.78	**	+	**	+
NT1RF2004750	8.14	5.64	6.27	13.53	14.23	13.30	8.05	8.74	9.81	**	+		
NT1RF2004755	11.30	7.99	8.26	16.42	20.16	17.92	10.59	13.63	13.47	**	+		
NT1RF2004767	6.21	2.89	4.95	9.44	8.05	8.14	4.7	6.19	4.36	*	+		
NT1RF2004768	9.61	3.95	2.60	2.99	2.03	1.57	2.24	1.57	1.49				
NT1RF2004775	2.25	2.07	1.48	4.36	5.01	5.07	4.16	3.75	3.44	**	+	**	+
NT1RF2004791	14.05	7.61	6.73	8.91	10.03	9.17	7.11	6.72	8.15				
NT1RF2004794	41.53	28.26	27.09	43.02	36.69	32.68	39.95	33.86	41.52				
NT1RF2004795	3.77	2.11	2.19	3.89	7.37	3.74	3.78	5.26	5.25		*	+	
NT1RF2004799	5.43	1.93	3.24	6.30	6.15	4.50	3.93	5.78	3.84				
NT1RF2004802	4.83	2.53	3.34	7.41	6.03	5.58	2.16	3.27	3.61	*	+		
NT1RF2004810	3.12	1.86	2.24	8.72	9.56	6.30	5.77	5.46	6.09	**	+	**	+
NT1RF2004816	4.85	3.14	2.65	6.62	9.96	5.26	6.09	3.65	4.78				
NT1RF2004837	13.44	8.28	7.12	11.51	16.25	16.53	19.77	16.72	17.56		*	+	
NT1RF2004841	2.64	1.81	1.21	3.03	4.37	3.11	1.94	3.01	1.95				
NT1RF2004847	16.48	11.83	12.45	15.24	18.08	16.57	16.4	14.80	14				
NT1RF2004861	1.52	1.27	1.44	3.27	3.09	3.21	1.26	1.81	1.52	**	+		
NT1RF2004897	1.25	0.88	1.99	3.40	2.11	1.91	1.21	2.22	1.75				
NT1RF2004932	10.00	7.17	11.03	13.12	14.42	13.51	9.72	9.64	9.65	*	+		
NT1RF2004933	1.78	1.31	1.88	3.51	3.60	2.84	3.51	3.18	3.33	**	+	**	+
NT1RF2004936	4.87	2.22	1.77	6.48	8.16	3.31	4.73	2.49	2.48				
NT1RF2004951	5.43	2.53	1.87	3.02	4.24	3.02	2.87	3.70	11.67				
NT1RF2004959	8.45	5.08	5.37	8.17	7.86	9.93	4.85	5.55	4.46				
NT1RF2004961	5.21	3.54	2.31	7.99	9.20	8.11	4.59	5.46	6.53	**	+		
NT1RF2004963	4.01	2.64	2.72	5.11	4.60	4.41	3.88	3.76	3.58	*	+		
NT1RF2004966	2.57	2.53	3.68	2.80	3.88	2.77	2.12	3.33	4.07				
NT1RF2004967	2.23	2.61	2.86	7.50	6.79	8.12	3.33	4.64	3.83	**	+	*	+
NT1RF2004974	1.95	1.80	1.93	2.56	3.12	2.39	3.76	2.94	0.71	*	+		
NT1RF2004978	6.88	2.95	2.57	5.63	7.09	3.07	4.98	3.62	3.21				
NT1RF2004982	1.90	1.58	1.47	6.52	6.96	3.08	1.22	2.05	1.93	*	+		
NT1RF2004985	24.53	11.76	13.37	30.81	35.00	31.74	21.76	19.69	22.43	*	+		
NT1RF2004999	4.87	3.06	2.28	6.14	7.08	4.89	3.19	4.04	3.16				
NT1RF2005000	3.68	2.30	2.22	2.75	3.93	3.69	1.87	3.37	3.49				
NT1RF2005001	3.57	1.78	2.11	2.93	4.06	3.83	3.59	4.32	2.86				
NT1RF2005003	4.87	3.07	2.71	7.63	8.71	7.19	5.69	5.10	6.02	**	+	*	+
NT1RF2005012	6.73	4.06	6.10	5.56	7.80	5.46	4.68	6.28	4.92				
NT1RF2005018	7.22	3.93	3.53	6.32	10.68	5.74	4.63	4.62	5.01				
NT1RF2005020	17.60	10.40	7.54	8.46	8.34	5.63	6.22	5.13	5.78				
NT1RF2005023	4.95	2.69	3.66	5.26	6.40	4.90	4.15	3.47	4.07				
NT1RF2005027	22.97	13.64	17.61	9.34	8.66	7.21	22.54	22.24	24.2	*	-		
NT1RF2005031	1.59	1.04	2.13	1.35	2.05	1.82	1.32	2.27	2.73				
NT1RF2005035	12.28	9.78	9.98	17.38	24.50	17.61	24.38	23.70	30.68	*	+	**	+
NT1RF2005037	3.95	3.48	2.80	4.77	7.93	4.42	2.77	3.83	4.79				
NT1RF2005038	1.07	1.01	1.27	2.71	2.99	1.89	1.22	3.15	1.8	*	+		
NT1RF2005048	8.09	4.51	4.12	7.64	8.60	7.45	7.59	5.31	4.79				
NT1RF2005069	25.41	8.17	11.97	37.61	33.07	31.21	30.69	32.10	38.73	*	+	*	+
NT1RF2005073	4.93	2.00	2.06	7.13	4.92	3.75	2.76	2.91	4.07				
NT1RF2005097	4.59	2.92	2.93	3.87	3.63	3.16	2.4	2.61	2.69				
NT1RF2005108	3.21	2.75	1.61	3.23	2.96	2.92	1.57	2.62	2.37				
NT1RF2005116	9.11	5.71	5.87	6.08	9.75	7.92	7.26	7.17	8.23				
NT1RF2005126	8.28	8.63	9.53	6.69	10.50	10.65	4.18	6.96	4.15	*	-		
NT1RF2005135	3.79	3.03	2.85	3.91	5.50	2.16	3.03	4.16	3.27				

92/180

【0733】

【表262】

1661

1662

NT2RP2005139	3.84	1.72	1.31	3.14	3.97	2.27	2.16	2.33	2.71										
NT2RP2005140	6.44	3.34	1.76	2.06	2.19	1.94	1.62	2.45	4.48										
NT2RP2005144	7.59	4.23	3.57	8.56	9.25	7.68	4.75	8.24	8.15										
NT2RP2005147	3.33	1.34	1.33	2.20	2.64	3.04	4.92	2.57	1.84										
NT2RP2005148	4.87	2.63	2.05	4.55	5.06	4.19	2.73	4.23	3.35										
NT2RP2005159	3.35	2.32	2.38	3.01	3.13	3.18	2.03	3.88	1.9										
NT2RP2005163	3.09	1.68	1.72	3.70	3.44	2.30	2.24	3.35	2.16										
NT2RP2005163	25.94	15.25	17.25	21.49	24.77	28.25	17.62	25.86	21.18										
NT2RP2005168	4.54	2.65	2.28	2.25	4.03	2.91	2.1	1.69	2.5										
NT2RP2005181	9.05	4.31	4.53	4.26	4.18	3.03	3.8	2.76	3.1										
NT2RP2005284	8.22	7.14	6.39	7.26	7.87	6.45	7	4.58	3.93										
NT2RP2005319	6.43	4.48	4.74	6.61	6.15	4.27	4.15	5.58	7.21										
NT2RP2005327	6.13	3.78	3.14	9.09	11.14	7.97	3.82	5.07	8.88	*	+								
NT2RP2005337	27.33	18.84	15.64	23.79	22.48	23.44	22.52	21.69	18.11										
NT2RP2005339	3.74	1.34	1.71	2.73	2.86	2.63	2.66	2.69	2.3										
NT2RP2005347	2.49	2.14	1.98	4.28	4.68	4.69	2.63	2.43	2.5	**	+								
NT2RP2005354	9.04	3.29	3.29	8.47	7.53	8.80	7.01	6.79	4.08										
NT2RP2005379	4.99	2.71	2.82	6.57	6.85	4.80	6.2	6.16	8.3			*	+						
NT2RP2005376	9.47	6.54	6.31	10.41	11.77	12.24	5.39	7.57	7.48	*	+								
NT2RP2005387	4.80	3.96	2.36	5.91	7.62	8.20	5.51	5.27	7.29	*	+								
NT2RP2005388	3.78	1.10	1.91	4.67	4.69	3.22	2.56	2.68	2.46										
NT2RP2005389	3.95	2.82	3.63	10.36	10.31	13.45	7.04	9.38	8.68	**	+	**	+						
NT2RP2005393	4.69	3.98	2.48	2.80	6.37	4.36	1.98	2.19	8.18										
NT2RP2005315	4.50	2.51	3.53	6.84	5.84	6.72	4.55	3.38	3.33	*	+								
NT2RP2005323	8.85	3.21	3.77	5.49	9.42	5.85	5.53	11.41	21.87										
NT2RP2005325	13.28	7.03	7.32	9.81	8.97	5.93	11.14	10.62	11.49										
NT2RP2005336	12.73	6.78	5.54	13.58	10.27	12.67	8.85	6.83	5.91										
NT2RP2005343	6.02	1.89	2.05	7.45	9.65	7.01	10.08	10.85	12.82	*	+	**	+						
NT2RP2005344	1.85	1.66	1.47	2.08	2.88	1.92	2.74	2.45	3.15			**	+						
NT2RP2005347	4.37	2.71	1.89	5.25	5.00	4.78	3.35	2.93	2.34										
NT2RP2005354	12.00	6.61	6.14	17.43	12.77	12.49	8.48	9.88	9.01										
NT2RP2005358	4.88	3.45	2.64	4.51	4.14	3.14	3.97	2.53	1.99										
NT2RP2005368	7.88	5.76	2.39	6.48	5.68	6.59	4.31	3.84	6.35										
NT2RP2005378	18.33	8.81	8.98	11.83	10.64	10.23	12.69	11.85	15.35										
NT2RP2005391	11.21	5.99	4.87	8.42	9.50	6.15	7.72	6.42	7.6										
NT2RP2005393	7.14	5.04	4.09	7.19	7.55	7.32	5.14	5.24	6.8										
NT2RP2005487	4.70	3.27	2.59	4.12	5.86	4.29	4.19	4.07	6.46										
NT2RP2005419	2.03	2.94	2.38	2.87	3.30	2.26	2.46	2.93	2.38										
NT2RP2005425	3.16	1.77	1.43	6.79	4.57	5.63	3.84	5.07	4.35	*	+	*	+						
NT2RP2005429	5.40	3.41	3.71	7.74	6.15	6.01	3.54	4.45	2.89	*	+								
NT2RP2005436	11.49	5.63	5.95	16.34	13.38	12.70	9.59	8.94	10.22	*	+								
NT2RP2005441	2.64	2.24	1.49	4.39	3.02	4.62	2.37	2.61	2.65	*	+								
NT2RP2005442	6.72	3.80	3.11	6.35	6.53	5.07	6.08	5.33	7.07										
NT2RP2005444	14.62	10.40	7.75	7.08	9.17	7.55	7.37	6.88	8.45										
NT2RP2005453	1.54	2.20	1.49	7.85	9.47	8.01	8.67	8.44	9.15	**	+	**	+						
NT2RP2005457	15.76	12.87	16.87	26.94	13.90	21.92	12.51	12.21	12.15										
NT2RP2005458	1.63	1.87	2.03	5.92	5.93	3.89	2.67	3.64	6.17	**	+								
NT2RP2005463	4.65	3.64	4.43	7.72	7.84	5.33	6.02	6.93	5.84	*	+	*	+						
NT2RP2005464	11.98	9.14	6.68	11.62	10.20	8.75	5.59	3.86	4.74			*	+						
NT2RP2005465	4.57	3.64	2.60	8.98	7.23	8.68	2.44	5.04	5.3	**	+								
NT2RP2005473	10.01	4.28	4.30	7.95	7.14	5.73	3.03	3.71	5.35										
NT2RP2005476	5.22	3.10	3.30	10.18	12.60	10.12	5.36	4.72	5.84	**	+								
NT2RP2005498	5.25	3.96	4.56	6.13	9.22	5.46	5.31	3.92	5.71										
NT2RP2005491	15.97	8.85	12.00	4.52	5.86	4.78	8.53	10.16	9.41	*	-								
NT2RP2005495	2.68	2.26	2.48	2.05	3.65	3.42	3.01	4.37	2.75										
NT2RP2005496	9.04	5.08	6.06	16.30	11.28	12.12	9.01	10.34	6.32	*	+								

93/180

【0734】

【表263】

1663

1664

NT2RF2005498	6.78	2.60	2.45	2.62	6.63	3.50	3.33	3.34	4.18										
NT2RF2005501	4.44	2.53	2.65	2.38	4.12	2.69	2.07	3.28	2.78										
NT2RF2005506	5.72	4.30	3.10	5.43	9.55	6.10	24.52	21.82	25.02			**	+						
NT2RF2005509	6.91	5.58	4.63	12.32	11.78	9.14	5.34	8.99	8.48	*	+								
NT2RF2005514	3.36	2.23	2.53	3.26	5.18	4.19	3.03	4.16	4.55	*	+								
NT2RF2005520	10.34	5.10	5.86	6.07	8.22	5.46	3.87	3.79	3.08										
NT2RF2005525	6.12	4.01	5.33	8.58	7.75	8.13	5.26	8.01	5.47	*	+								
NT2RF2005531	0.65	1.10	1.57	2.53	1.56	1.74	1.49	2.39	1.21										
NT2RF2005535	36.57	17.31	21.13	93.90	73.03	67.87	27.53	17.14	25.99	**	+								
NT2RF2005539	10.87	6.53	4.81	8.43	9.17	6.85	6.76	6.87	5.25										
NT2RF2005540	2.81	2.63	2.81	7.15	6.27	3.67	4.42	5.46	9.74	**	+								
NT2RF2005541	5.40	3.42	2.70	8.82	9.81	10.04	7.49	7.37	5.44	**	+	*	+						
NT2RF2005549	3.91	1.98	1.81	3.23	5.51	2.41	2.43	3.46	2.97										
NT2RF2005553	3.52	2.33	3.66	6.38	7.53	5.49	7.54	10.56	6.47	*	+	*	+						
NT2RF2005557	7.00	5.12	11.72	16.35	11.47	12.41	6.34	5.80	8.04										
NT2RF2005561	5.51	4.09	4.45	13.70	13.23	10.54	6.26	5.62	5.86	**	+								
NT2RF2005586	7.40	3.49	4.35	2.55	4.08	2.63	1.67	2.60	2.43										
NT2RF2005597	6.16	4.97	3.02	4.57	4.34	4.57	4.67	4.40	3.08										
NT2RF2005600	4.06	2.52	2.53	3.83	4.26	3.10	2.47	4.00	2.95										
NT2RF2005605	13.12	8.01	6.74	12.67	14.30	12.26	6.96	7.51	8										
NT2RF2005614	9.18	5.27	8.25	16.39	16.00	13.57	10.11	8.70	9.2	**	+								
NT2RF2005628	4.07	2.65	2.40	3.99	3.40	3.40	2.45	3.61	2.26										
NT2RF2005632	9.20	6.36	7.23	6.07	7.94	5.76	4.64	4.67	6.34										
NT2RF2005633	3.64	3.42	2.57	5.77	4.33	3.82	2.82	3.85	3.3										
NT2RF2005635	3.95	2.73	2.06	3.40	4.38	2.94	2.4	2.42	3.18										
NT2RF2005637	2.20	1.05	1.68	13.21	4.02	4.55	2.2	2.55	5.6										
NT2RF2005640	3.47	1.55	1.53	2.16	1.23	2.22	1.96	2.66	2.84										
NT2RF2005645	6.42	3.67	2.99	5.68	11.68	7.34	5.29	6.74	5.73										
NT2RF2005651	4.09	3.02	3.19	6.89	11.77	5.52	3.81	4.33	6.7										
NT2RF2005654	5.50	3.61	4.20	6.10	7.84	5.96	4.19	5.64	4.96										
NT2RF2005666	4.54	3.08	3.45	5.18	6.63	4.14	4.25	3.69	7.2										
NT2RF2005669	6.09	5.35	5.64	8.34	9.73	9.01	4.66	6.00	6.82	**	+								
NT2RF2005670	2.87	2.57	1.87	5.75	5.68	2.97	1.68	2.33	3.03										
NT2RF2005671	10.41	3.42	4.33	5.10	6.52	3.51	3.46	4.47	6.12										
NT2RF2005675	11.31	4.30	4.30	8.54	8.22	4.79	7.64	6.94	9.43										
NT2RF2005683	9.32	5.43	5.87	8.08	9.48	5.92	5.85	4.94	4.56										
NT2RF2005689	3.18	1.30	1.52	3.24	4.46	3.75	2.33	3.71	3.54										
NT2RF2005694	4.53	2.30	2.18	4.82	3.54	4.62	3.22	3.77	3.78										
NT2RF2005701	22.21	13.84	17.86	22.12	25.56	24.08	18.18	17.70	22.41										
NT2RF2005712	2.84	3.06	3.02	3.90	3.94	3.10	1.15	2.49	1.88			*	-						
NT2RF2005719	2.26	1.27	0.72	3.09	3.04	2.67	2.23	1.46	2.56	*	+								
NT2RF2005722	11.76	8.52	5.52	18.21	24.59	18.10	8.26	9.21	12.37	*	+								
NT2RF2005723	4.68	2.75	2.29	7.35	6.52	3.86	4.39	4.70	2.79										
NT2RF2005726	5.41	2.39	2.73	5.77	4.51	4.16	3.27	4.19	3.67										
NT2RF2005729	5.30	2.58	2.08	6.82	6.27	4.01	3.21	5.54	3.89										
NT2RF2005731	0.50	0.60	0.63	1.06	1.43	0.80	0.71	2.81	0.87	*	+								
NT2RF2005732	8.98	3.61	4.01	6.71	6.46	5.79	4.23	7.06	7.16										
NT2RF2005737	10.83	8.16	10.12	14.63	17.80	12.60	12.9	11.51	9.06	*	+								
NT2RF2005741	5.83	2.63	2.65	3.36	3.80	2.41	3.96	2.72	3.47										
NT2RF2005748	3.52	1.63	2.33	2.18	2.64	1.48	3.11	2.62	2.38										
NT2RF2005752	5.37	3.43	3.73	6.46	5.65	5.66	6.55	3.67	3.82										
NT2RF2005753	22.04	14.07	18.05	15.96	24.14	20.59	21.63	18.25	19.82										
NT2RF2005763	6.73	2.47	2.52	3.25	3.61	3.70	1.84	3.88	3.22										
NT2RF2005767	2.43	2.60	2.16	6.91	6.56	7.20	3.36	3.03	4.72	**	+	*	+						
NT2RF2005773	15.62	10.12	12.99	19.66	19.02	17.26	17.15	13.07	15.8	*	+								
NT2RF2005774	10.33	5.72	6.91	21.21	24.60	21.03	9.42	7.55	8.22	**	+								

94/180

【0735】

【表264】

1665

1666

NT2RP2005775	4.39	1.98	1.42	2.12	2.56	2.56	2.19	2.08	1.67										
NT2RP2005781	5.85	3.98	3.29	6.76	5.57	5.04	4.75	3.50	4.17										
NT2RP2005784	11.14	6.72	5.29	8.15	8.38	8.40	7.85	8.40	10.24										
NT2RP2005789	4.85	3.33	3.28	5.63	7.04	4.46	3.88	3.70	4.09										
NT2RP2005799	1.71	1.81	1.37	3.76	5.36	2.16	2.16	2.19	2.43			*							
NT2RP2005804	6.19	3.18	3.30	4.57	7.49	6.42	5.55	5.88	4.72										
NT2RP2005812	3.92	3.04	2.54	4.78	6.17	3.21	2.98	4.18	4.04										
NT2RP2005815	2.54	2.17	3.20	3.81	3.69	2.58	2.35	2.98	1.88										
NT2RP2005835	14.04	7.44	6.79	14.50	10.00	10.84	9.86	7.11	11.61										
NT2RP2005841	6.35	3.23	3.13	5.70	4.93	4.82	5.84	3.68	4.77										
NT2RP2005853	3.23	3.29	2.96	6.28	6.53	5.74	4.87	4.09	5.28	**		+	*						
NT2RP2005857	8.95	4.28	4.74	6.65	7.52	6.19	1.63	2.12	1.8										
NT2RP2005859	5.38	4.41	5.54	4.28	5.42	3.86	2.87	3.84	3.87			*							
NT2RP2005869	3.02	1.60	2.64	2.92	4.01	2.37	2.32	4.74	1.81										
NT2RP2005863	4.66	2.88	2.88	3.96	3.85	3.93	2.02	2.05	1.69										
NT2RP2005868	3.44	1.57	1.65	4.52	4.28	2.97	2.38	3.85	2.89										
NT2RP2005876	13.61	7.01	5.40	17.03	13.16	6.91	8.8	8.61	107										
NT2RP2005878	6.92	4.37	4.13	11.06	12.33	11.73	5.81	7.81	6.82	**		+							
NT2RP2005883	1.59	1.56	1.08	3.31	2.84	2.42	3.91	4.53	4.86	**		+	**						
NT2RP2005886	8.60	4.98	6.40	10.11	11.16	11.42	6.19	6.08	5.43	*		+							
NT2RP2005887	5.47	3.26	3.97	12.05	12.81	9.32	13.75	10.35	15.02	**		+	**						
NT2RP2005890	7.74	6.08	7.50	6.23	6.35	4.71	2.57	2.56	1.86			**							
NT2RP2005891	3.39	2.76	2.57	3.81	4.07	4.20	2.43	3.04	3.13	*		+	*						
NT2RP2005902	1.86	0.89	1.33	3.39	3.77	2.15	2.13	2.79	3.13	*		+	*						
NT2RP2005908	9.46	5.71	4.03	9.28	7.93	10.45	6.03	6.26	6.92										
NT2RP2005917	7.43	5.84	5.10	9.51	9.65	7.14	3.72	5.75	4.41										
NT2RP2005933	6.32	4.20	3.63	5.57	7.02	4.50	3.29	2.73	4.08										
NT2RP2005941	9.03	6.94	7.01	7.65	13.07	8.78	10.41	9.47	5.87										
NT2RP2005942	3.02	2.03	1.79	3.90	4.09	3.96	2.56	2.66	2	*		+							
NT2RP2005946	6.57	4.95	5.93	3.90	3.86	3.37	2.5	2.94	2.41	*		-	**						
NT2RP2005979	12.30	10.25	11.94	15.87	16.05	15.06	14.9	13.37	14.97	**		+	*						
NT2RP2005980	3.71	2.65	2.25	7.90	7.37	4.49	4.13	4.23	2.71	*		+							
NT2RP2005994	5.01	2.60	2.01	2.75	4.22	1.07	2.23	3.11	2.43										
NT2RP2006004	2.32	1.82	1.35	2.43	4.21	2.56	2.36	3.37	2.03										
NT2RP2006013	4.44	2.15	4.45	6.09	6.99	3.28	4.68	5.22	4.41										
NT2RP2006023	21.60	12.40	20.04	37.44	49.33	45.44	22.61	22.79	24.39	**		+							
NT2RP2006028	5.34	3.20	3.73	4.07	4.23	2.81	3.39	4.81	5.42										
NT2RP2006038	0.34	0.06	1.28	0.43	0.83	3.61	0.25	1.80	0.18										
NT2RP2006042	8.65	5.14	6.93	7.32	7.79	6.34	7.56	7.82	9.4										
NT2RP2006049	5.05	2.75	2.80	12.32	12.87	10.73	8.05	8.08	7.82	**		+	**						
NT2RP2006062	2.31	2.64	1.44	1.42	2.55	2.98	1.26	2.10	2.6										
NT2RP2006067	3.69	1.67	1.24	3.57	3.44	2.48	2.2	3.85	3.27										
NT2RP2006064	12.49	6.77	9.83	12.13	10.85	6.00	10.28	6.81	5.57										
NT2RP2006068	3.25	3.63	2.31	8.60	6.86	6.64	4.6	5.09	2.54	**		+							
NT2RP2006069	1.08	0.69	0.92	0.88	1.74	0.95	0.92	1.48	1.42										
NT2RP2006071	2.73	3.23	2.31	5.07	7.66	5.45	2.92	4.00	2.55	*		+							
NT2RP2006079	3.70	1.69	2.79	3.57	5.20	3.82	3.74	3.63	2.49										
NT2RP2006092	3.65	2.47	2.47	3.19	3.41	3.44	2.36	2.80	2.77										
NT2RP2006097	24.23	9.76	10.66	21.53	18.65	14.12	10.2	9.65	12.96										
NT2RP2006098	4.17	2.27	1.77	4.26	4.04	2.86	4.03	6.51	3.5										
NT2RP2006099	4.48	2.99	2.12	5.82	5.86	5.18	3.32	4.84	3.86	*		+							
NT2RP2006100	3.88	1.55	1.83	3.98	4.92	3.85	2.1	3.68	2.65										
NT2RP2006103	10.54	3.86	5.78	3.37	2.55	1.88	2.21	3.36	1.85										
NT2RP2006106	8.45	4.11	4.04	6.45	5.69	6.30	4.48	6.29	4.22										
NT2RP2006127	9.00	6.34	7.56	9.10	8.66	7.24	9.22	10.10	8.49										
NT2RP2006134	1.55	1.02	1.47	1.76	1.82	1.93	1.55	2.52	1.39	*		+							

95/180

【0736】

【表265】

1667

1668

NT2RF2006141	5.76	3.11	3.17	3.84	5.50	4.54	3.67	3.75	3.95										
NT2RF2006146	7.93	5.66	5.17	12.63	13.99	9.56	6.76	6.08	6.36	*	+								
NT2RF2006176	4.45	2.26	1.67	6.40	4.88	5.22	2.44	3.34	5.68	*	+								
NT2RF2006181	1.58	1.06	1.00	1.37	3.24	3.22	1.23	2.94	1.73										
NT2RF2006184	23.94	15.54	16.09	22.96	21.00	23.09	17.11	19.55	14.56										
NT2RF2006186	1.68	1.14	2.35	2.02	3.74	1.74	1.23	3.31	1.82										
NT2RF2006196	4.74	3.02	3.70	6.83	6.02	5.77	4.04	5.17	3.91	*	+								
NT2RF2006199	2.29	2.59	2.52	3.33	3.50	4.30	2.88	2.76	2.12	*	+								
NT2RF2006200	4.29	2.63	1.43	3.59	5.59	2.06	3.12	2.50	2.5										
NT2RF2006210	59.40	41.07	36.68	33.45	39.27	22.58	11.72	8.31	9.47			**	-						
NT2RF2006219	3.75	1.76	1.64	3.39	3.29	2.82	2.17	1.88	4.22										
NT2RF2006224	5.72	3.72	4.01	5.11	6.26	6.39	3.82	3.77	4.2										
NT2RF2006237	5.09	3.91	5.00	9.00	7.92	10.28	5.01	5.76	5.27	**	+								
NT2RF2006239	3.42	2.16	1.78	4.42	4.29	2.44	2.31	3.01	1.89										
NT2RF2006238	9.12	5.55	6.28	6.08	7.62	7.68	6.35	6.07	3.93										
NT2RF2006261	1.75	2.42	1.14	2.06	2.49	1.87	1.21	1.75	2.67										
NT2RF2006269	23.86	9.30	9.53	15.39	18.13	13.53	12.46	10.61	15.67										
NT2RF2006275	4.68	2.71	2.12	3.99	3.20	2.45	2.89	2.81	3.34										
NT2RF2006282	7.12	3.89	6.34	8.17	11.45	9.25	4.48	4.87	2.85	*	+								
NT2RF2006302	4.86	2.69	3.31	10.51	10.47	9.06	10	9.83	3.76	**	+								
NT2RF2006312	8.45	5.62	5.99	10.60	10.03	9.84	7.18	6.51	5.02	*	+								
NT2RF2006320	3.62	2.45	1.39	4.62	5.47	5.86	2.21	4.05	3.23	*	+								
NT2RF2006321	1.99	1.78	2.42	3.22	4.24	2.52	1.97	3.17	2.07										
NT2RF2006323	1.30	0.75	0.38	1.35	1.65	0.69	0.19	2.09	2.6										
NT2RF2006333	2.18	0.70	0.66	2.51	1.88	1.17	1.49	1.76	2.35										
NT2RF2006334	3.73	1.40	1.47	2.69	3.03	2.34	0.81	2.29	2.95										
NT2RF2006338	2.65	1.82	1.03	3.45	4.02	2.81	1.6	3.69	2.93										
NT2RF2006339	2.37	1.54	1.37	3.09	2.39	1.47	1.2	2.47	2.21										
NT2RF2006355	1.01	0.99	0.71	2.16	2.25	1.72	1.94	2.95	0.87	**	+								
NT2RF2006365	1.51	1.66	1.15	3.16	4.39	3.70	1.83	4.13	1.9	**	+								
NT2RF2006374	16.70	8.19	7.22	17.36	18.00	12.60	10.86	13.62	9.02										
NT2RF2006393	4.85	2.17	2.52	8.54	10.40	8.85	5.98	6.32	6.15	**	+	*	+						
NT2RF2006394	2.02	1.64	1.69	3.46	1.86	1.52	3.53	1.56	2.54										
NT2RF2006400	1.99	1.74	1.43	2.29	2.67	2.67	2.79	1.32	1.33	*	+								
NT2RF2006411	36.13	23.40	20.23	18.85	35.68	22.21	26.26	22.92	21.44										
NT2RF2006429	3.49	1.96	1.56	18.22	22.80	21.81	7.61	6.72	8.71	**	+	**	+						
NT2RF2006435	2.88	2.61	2.07	4.19	4.16	3.86	3.51	3.02	3.91	**	+	*	+						
NT2RF2006436	4.50	2.57	2.37	11.47	10.14	10.45	15.06	14.83	12.68	**	+	**	+						
NT2RF2006441	5.48	3.11	4.57	12.23	11.44	10.95	9.38	9.44	9.01	**	+	**	+						
NT2RF2006447	3.63	2.74	2.87	7.53	5.11	1.37	2.09	1.56	0.94			*	-						
NT2RF2006454	3.45	1.48	1.32	2.04	2.21	2.24	3.02	1.84	0.51										
NT2RF2006455	3.08	1.02	1.42	3.46	1.52	2.11	2.25	1.46	1.25										
NT2RF2006456	3.43	1.56	1.38	1.87	3.29	2.20	1.39	3.00	3.52										
NT2RF2006464	7.78	4.58	3.90	5.55	4.82	4.88	3.6	3.54	5.67										
NT2RF2006467	5.66	2.72	2.67	10.90	8.83	10.03	7.29	6.50	9.64	**	+	*	+						
NT2RF2006472	7.44	3.78	3.97	8.69	8.19	8.22	5	4.65	10.62										
NT2RF2006474	8.86	5.98	7.97	27.71	30.65	24.91	30	37.03	33.44	**	+	**	+						
NT2RF2006475	5.74	3.11	2.17	15.80	11.04	13.89	8.72	6.46	9.93	**	+	*	+						
NT2RF2006476	14.81	5.52	5.85	6.07	6.15	5.06	6.6	4.52	6.65										
NT2RF2006501	10.57	4.49	3.64	10.98	10.15	9.25	4.35	4.19	5.76										
NT2RF2006512	10.18	4.42	5.26	7.98	9.45	6.77	5.81	5.75	6.47										
NT2RF2006526	2.38	0.63	1.13	1.33	3.17	1.44	1.31	2.50	2.57										
NT2RF2006527	6.04	4.50	5.90	6.98	6.77	8.30	6.37	6.06	6.05										
NT2RF2006534	1.08	0.58	0.52	1.10	1.90	2.81	1.51	1.54	1.55			*	+						
NT2RF2006537	7.96	4.17	4.11	12.78	11.80	12.98	5.84	7.01	9.26	**	+								
NT2RF2006543	2.53	2.49	1.25	3.82	3.98	2.63	5.74	3.55	4.9			*	+						

96/180

[0737]

[表266]

1669

1670

NT2RF2006554	2.93	1.44	1.64	4.14	5.11	5.65	3.05	2.87	4.34	**	+		
NT2RF2006565	2.42	3.04	1.97	5.84	7.27	4.73	5.76	4.50	8.32	*	+	*	
NT2RF2006571	15.53	8.80	8.87	9.19	10.25	5.31	9.49	9.09	15.1				
NT2RF2006573	3.03	1.23	1.11	3.74	3.96	3.02	2.6	2.13	2.11				
NT2RF2006596	5.73	3.98	4.61	7.93	8.72	6.43	5.28	3.71	6.12	*	+		
NT2RF2006601	37.52	34.93	32.64	41.04	41.47	32.68	27.39	28.66	36.43				
NT2RF3000002	3.95	2.25	3.29	4.37	7.61	7.60	3.47	4.83	7.96	*	+		
NT2RF3000011	4.07	2.64	1.62	5.92	4.70	5.14	3.96	3.12	4.24	*	+		
NT2RF3000014	3.17	3.00	2.39	9.14	11.05	8.39	7.15	7.48	8.57	**	+	**	+
NT2RF3000016	9.66	5.49	5.68	6.73	6.36	7.49	4.75	5.66	6.35				
NT2RF3000022	4.96	2.03	2.47	3.53	3.43	2.45	3.24	3.89	7.1				
NT2RF3000024	12.74	9.32	13.69	28.77	37.69	22.23	11.49	12.80	14.79	*	+		
NT2RF3000031	4.64	2.28	2.98	4.90	4.09	5.50	4.12	3.94	3.26				
NT2RF3000034	4.51	3.69	3.49	3.95	4.58	4.75	3.38	3.05	3.23				
NT2RF3000037	15.49	9.32	10.69	13.56	14.15	12.81	7.78	9.45	8.16				
NT2RF3000040	2.98	2.45	1.73	1.43	1.95	2.12	0.99	2.09	2.1				
NT2RF3000041	10.75	6.47	4.78	19.57	16.79	13.38	9.67	7.12	9.17	*	+		
NT2RF3000046	5.16	2.85	2.89	6.40	9.13	5.39	4.23	3.75	6.16				
NT2RF3000047	6.44	3.75	3.07	4.50	4.32	4.37	3.44	4.24	4.69				
NT2RF3000049	3.94	3.36	1.85	3.67	6.35	6.22	5.02	4.43	8.2				
NT2RF3000059	7.94	4.67	6.52	13.03	15.60	12.76	7.92	7.66	10.86	**	+		
NT2RF3000061	6.26	3.23	4.99	9.29	9.59	8.78	5.46	7.17	6.65	**	+		
NT2RF3000064	6.09	3.47	4.38	5.67	6.99	5.26	5.01	4.84	5.62				
NT2RF3000065	3.24	2.73	0.81	4.89	4.66	2.53	2.67	2.43	3.79				
NT2RF3000066	2.70	3.24	1.60	2.60	3.66	2.74	3.75	2.94	3.3				
NT2RF3000069	4.21	2.87	2.12	3.45	3.50	3.02	3.35	3.22	4.21				
NT2RF3000063	7.78	5.44	6.74	6.64	5.14	7.47	6.5	8.34	4.12				
NT2RF3000068	1.30	1.86	2.21	1.64	3.20	2.26	2.1	3.07	3.12				
NT2RF3000069	3.21	2.16	2.26	10.79	10.68	7.75	8.64	7.90	7.98	**	+	**	+
NT2RF3000072	2.08	1.15	1.36	3.34	2.75	2.73	2.05	3.07	2.12	*	+		
NT2RF3000080	12.90	8.84	11.62	14.83	16.14	12.41	14.4	11.56	12.15				
NT2RF3000085	4.82	2.44	2.00	2.73	3.07	3.01	2.95	2.26	2.49				
NT2RF3000087	12.35	7.36	5.97	19.26	20.25	18.12	12.89	8.99	10.11	**	+		
NT2RF3000092	2.83	2.11	1.59	4.04	2.45	1.56	2.71	2.87	2.87				
NT2RF3000109	1.75	1.89	2.71	5.02	4.39	3.90	1.58	3.14	1.97	**	+		
NT2RF3000119	10.48	4.74	6.30	7.48	8.15	6.85	5.44	7.67	7.52				
NT2RF3000125	9.53	6.24	6.75	10.54	13.59	12.33	7.17	8.82	6.83	*	+		
NT2RF3000131	13.37	7.84	8.67	12.43	13.75	13.12	11.27	10.91	10.26				
NT2RF3000134	8.39	4.00	4.04	11.86	8.47	11.09	6.37	5.88	5.18	*	+		
NT2RF3000137	7.33	3.86	4.11	4.55	10.23	5.70	5.38	4.77	5.25				
NT2RF3000143	8.58	2.85	3.30	8.25	6.01	4.98	4.9	4.68	4.31				
NT2RF3000148	6.50	3.03	2.82	4.77	5.93	4.35	4.39	4.87	3.31				
NT2RF3000149	7.40	4.34	3.38	4.95	6.06	4.71	3.63	4.88	5.43				
NT2RF3000163	5.34	2.10	2.73	5.49	7.84	4.53	2.61	3.70	2.68				
NT2RF3000164	17.73	9.34	8.35	13.43	12.52	14.26	15.5	18.99	23.37				
NT2RF3000169	2.79	1.47	1.93	3.28	2.66	2.80	2.69	4.02	3.92				
NT2RF3000171	30.99	20.17	24.95	41.61	37.53	33.55	22.47	25.44	33.88	*	+		
NT2RF3000172	5.29	2.13	2.18	3.70	4.85	1.88	2.31	1.91	2.23				
NT2RF3000186	16.37	8.43	6.94	11.35	12.10	6.88	5.69	5.57	6.97				
NT2RF3000197	2.96	2.49	2.66	5.21	6.67	3.78	2.54	2.96	3.64	*	+		
NT2RF3000201	11.54	5.67	6.73	11.59	11.99	10.04	5.11	5.52	10.33				
NT2RF3000204	3.53	2.05	1.72	2.68	3.65	3.34	1.75	3.41	1.98				
NT2RF3000207	4.88	2.36	2.46	3.16	3.56	3.29	4.1	5.13	6.04				
NT2RF3000216	8.62	6.38	5.44	6.42	8.59	6.54	8.63	6.16	7.72				
NT2RF3000229	2.88	1.23	2.50	2.75	3.70	2.71	2.57	3.23	2.33				
NT2RF3000231	4.47	2.97	2.52	4.75	5.82	4.37	3.89	3.79	4.34				

97/180

【0738】

【表267】

1671

1672

NT2RF3000233	7.80	2.59	4.87	14.07	13.78	10.58	4.43	6.16	6.42	*	+		
NT2RF3000233	4.29	2.04	3.30	4.16	4.02	3.58	3.88	4.05	3.95				
NT2RF3000234	5.82	3.69	3.99	6.88	6.24	5.76	5.09	5.25	5.52				
NT2RF3000235	4.07	2.16	2.75	4.46	3.39	3.79	3.35	4.56	3.36				
NT2RF3000239	7.80	3.65	4.61	5.36	6.98	5.05	3.92	4.89	7.01				
NT2RF3000247	2.30	1.21	1.95	2.12	2.01	2.94	1.85	3.86	2.35				
NT2RF3000251	8.89	5.54	6.24	11.87	10.35	8.87	9.19	9.33	7.77				
NT2RF3000253	15.04	4.46	4.08	9.00	9.21	5.83	5.52	5.10	6.74				
NT2RF3000255	5.13	2.85	2.23	3.53	3.93	3.32	2.12	2.52	4.46				
NT2RF3000263	7.20	3.34	3.67	7.23	8.28	5.67	4.7	4.54	3.81				
NT2RF3000266	13.99	6.47	5.93	16.36	16.88	13.38	9.91	14.29	13.15				
NT2RF3000267	4.19	1.73	1.51	3.08	4.17	2.39	2.28	3.19	2.31				
NT2RF3000271	7.47	3.16	2.85	7.84	6.39	5.57	3.5	5.30	3.78				
NT2RF3000278	3.14	2.04	3.02	4.79	6.42	5.26	3.22	5.70	4.21	**	+		
NT2RF3000281	7.14	3.51	4.30	9.39	7.57	6.94	6.62	8.48	7.76				
NT2RF3000292	2.43	1.31	1.46	1.66	2.08	1.80	2.97	2.36	1.82				
NT2RF3000299	3.32	1.72	2.64	3.50	2.85	1.65	3.49	2.65	2.85				
NT2RF3000304	7.20	4.06	3.87	3.27	5.90	6.50	4.23	4.68	5.46				
NT2RF3000310	9.88	5.44	4.97	10.57	8.79	8.65	8.38	7.53	9.91				
NT2RF3000312	4.71	2.11	3.36	4.19	4.91	4.91	2.11	3.53	4.02				
NT2RF3000326	9.82	2.79	5.46	8.18	6.79	9.80	7.95	7.10	16.94				
NT2RF3000332	30.65	18.22	26.99	58.85	49.93	31.40	36.14	39.97	34.74		*	+	
NT2RF3000334	2.18	1.49	1.41	2.10	2.20	2.50	2.87	1.62	1.63				
NT2RF3000326	4.07	2.09	2.65	6.40	4.79	6.20	5.05	3.50	3.68	*	+		
NT2RF3000329	8.08	3.03	2.39	13.04	10.42	8.93	5.43	5.08	6.48	*	+		
NT2RF3000336	6.13	3.81	4.47	3.99	4.93	3.61	5.76	6.52	5.37				
NT2RF3000333	3.58	1.99	1.19	2.09	2.88	2.04	2.14	2.57	2.31				
NT2RF3000341	13.34	6.74	7.40	16.98	14.13	16.48	11.16	11.51	12.58	*	+		
NT2RF3000344	2.19	2.15	1.77	2.27	1.91	1.50	1.56	1.76	2.32				
NT2RF3000345	0.88	0.64	0.51	3.07	2.22	3.27	0.95	0.77	2.11	**	+		
NT2RF3000348	112.18	53.12	48.19	87.36	67.82	76.37	170.4	141.05	175.2		*	+	
NT2RF3000358	13.69	7.30	6.99	9.25	9.00	7.77	7.42	5.74	8.01				
NT2RF3000359	10.64	6.49	5.35	19.00	17.38	16.68	15.5	13.49	16.08	**	+	*	+
NT2RF3000361	10.35	4.92	4.34	11.24	6.97	7.55	6.16	6.69	7.28				
NT2RF3000366	7.65	3.30	4.82	9.45	14.23	10.18	10.84	11.42	12.66	*	+	**	+
NT2RF3000378	4.91	3.67	4.88	5.34	6.49	6.00	4.34	4.99	3.64				
NT2RF3000384	6.56	5.43	5.50	8.93	9.13	11.76	6.91	6.90	7.16	*	+	*	+
NT2RF3000389	14.26	10.15	11.05	22.04	27.40	18.38	12.47	13.44	23.39	*	+		
NT2RF3000393	5.27	3.15	2.77	4.98	4.37	4.43	4.32	3.00	3.71				
NT2RF3000395	121.26	84.54	65.25	98.14	119.90	103.24	32.56	26.84	40.17		*		
NT2RF3000397	3.69	4.24	2.44	2.76	4.13	3.97	3.48	2.62	4.13				
NT2RF3000398	6.97	4.09	4.94	8.35	10.97	6.66	5.51	6.21	5.86				
NT2RF3000403	4.82	3.83	4.35	9.87	12.59	8.19	6.65	6.56	8.79	*	+	*	+
NT2RF3000418	4.00	2.62	2.61	8.58	12.65	8.62	5.36	6.28	7.5	**	+	*	+
NT2RF3000424	5.08	4.11	3.95	14.10	16.88	10.90	8.47	7.77	7.95	**	+	**	+
NT2RF3000427	2.50	1.80	2.77	5.73	6.63	8.27	3.99	5.02	3.87	**	+	*	+
NT2RF3000431	3.51	2.32	1.35	4.97	4.03	2.77	4.39	4.52	3.47				
NT2RF3000433	4.48	3.35	3.32	4.96	5.89	5.97	3.9	4.05	4.56	*	+		
NT2RF3000436	11.10	6.79	5.78	9.34	10.99	9.24	10.36	9.52	16.87				
NT2RF3000439	5.21	2.28	3.00	3.90	7.56	3.69	3.69	4.00	3.42				
NT2RF3000441	1.19	0.92	0.83	1.64	2.07	1.50	2.8	3.37	2.81	*	+	**	+
NT2RF3000444	2.26	2.00	1.85	2.13	2.91	3.48	2.82	2.26	2.53				
NT2RF3000448	3.48	2.24	3.61	8.12	11.89	8.40	5.13	4.03	6.51	**	+		
NT2RF3000449	5.49	2.45	3.20	2.67	4.04	3.28	1.61	2.66	1.96				
NT2RF3000451	5.47	3.68	2.74	2.86	3.50	4.17	4.01	4.24	4.31				
NT2RF3000456	4.82	4.21	3.70	3.94	5.59	4.41	3.70	5.42					

1673

1674

NT2RF3000469	6.78	3.61	3.73	8.08	8.65	7.41	13.4	12.29	10.37	*	+	**	+
NT2RF3000471	6.95	4.34	4.50	7.79	8.60	6.26	4.55	7.24	5.12				
NT2RF3000477	21.65	12.36	9.87	23.85	19.48	15.72	11.17	14.00	11.16				
NT2RF3000478	7.29	4.54	5.34	13.47	15.94	11.73	8.07	4.43	7.92	**	+		
NT2RF3000481	0.63	0.59	0.73	1.35	1.25	1.38	0.46	2.40	1.02	*	+		
NT2RF3000484	1.55	0.72	1.25	1.68	2.10	2.87	1.12	2.90	1.09				
NT2RF3000487	5.07	1.99	2.06	3.79	5.91	4.35	2.41	2.16	2.61				
NT2RF3000512	6.71	4.34	3.46	3.23	5.10	5.08	2.77	4.20	4.93				
NT2RF3000523	27.58	15.65	17.30	17.42	22.63	15.01	11.77	10.31	9.03				
NT2RF3000526	2.57	1.90	3.01	5.30	4.16	4.98	2.88	5.37	3.11	**	+		
NT2RF3000527	3.80	1.53	2.25	4.05	4.14	5.85	2.46	3.30	2.31				
NT2RF3000531	15.89	10.13	8.97	23.60	23.41	21.43	13.33	15.19	15.55	**	+		
NT2RF3000532	6.87	3.91	4.69	7.54	6.97	6.82	3.54	4.64	3.97				
NT2RF3000542	4.26	2.58	3.40	6.33	6.95	7.50	5.58	5.25	4.09	**	+		
NT2RF3000544	21.26	8.36	10.64	9.79	12.63	8.67	7.85	5.66	7.16				
NT2RF3000561	1.72	1.29	0.49	4.36	4.39	2.75	5.41	6.15	4.61	*	+	**	+
NT2RF3000562	5.35	3.52	2.70	6.24	5.67	6.85	4.69	5.36	4.51	*	+		
NT2RF3000572	2.48	1.13	0.91	1.41	2.33	1.29	1.51	2.72	1.83				
NT2RF3000582	2.70	1.06	2.14	1.55	1.76	2.00	1.13	2.91	1.43				
NT2RF3000584	3.87	1.71	2.00	3.83	3.38	4.15	1.95	3.50	3.43				
NT2RF3000586	4.68	3.18	3.48	5.21	5.82	4.88	4.06	4.66	4.73	*	+		
NT2RF3000590	3.21	1.61	2.30	2.02	1.87	2.52	1.95	2.50	2.25				
NT2RF3000592	2.67	1.26	1.45	1.25	2.76	1.46	1.33	1.90	1.13				
NT2RF3000596	20.65	9.80	8.82	23.94	26.59	16.13	11.86	9.91	14.07				
NT2RF3000599	3.31	1.61	2.33	3.96	4.14	2.63	2.43	4.34	3.3				
NT2RF3000603	4.81	2.59	2.37	5.30	5.93	6.54	3.73	4.56	4.65	*	+		
NT2RF3000605	2.51	1.85	1.50	3.30	3.59	2.96	2.17	4.09	3.29	*	+		
NT2RF3000607	7.51	5.55	8.79	5.67	5.09	3.67	3.76	3.78	3.57			*	-
NT2RF3000616	2.94	0.94	1.60	3.25	4.41	3.35	2.18	3.01	2.34				
NT2RF3000621	4.36	2.30	3.65	4.44	7.67	4.30	4.7	5.31	5.41				
NT2RF3000623	6.01	4.28	3.80	5.09	7.11	5.45	5.08	3.73	3.94				
NT2RF3000624	7.72	5.67	3.32	6.67	8.14	5.52	5.24	3.13	5.14				
NT2RF3000628	7.54	4.50	3.20	10.58	21.80	10.94	10.01	5.74	10.27				
NT2RF3000631	16.09	7.17	9.25	14.57	17.16	15.18	7.31	8.71	8.97				
NT2RF3000632	7.31	3.75	5.02	6.89	10.18	9.21	4.07	4.61	4.79				
NT2RF3000638	7.68	5.11	4.32	4.07	4.85	4.59	5.9	6.86	5.24				
NT2RF3000644	19.00	10.57	14.03	22.53	22.68	23.63	19.34	20.74	17.56	*	+		
NT2RF3000645	22.63	12.76	16.07	25.22	24.49	30.53	19.65	22.44	19.81				
NT2RF3000652	25.30	13.23	15.28	45.18	43.44	33.63	16.17	15.59	14.77	*	+		
NT2RF3000658	10.87	4.38	5.61	9.08	8.70	4.57	4.84	5.59	6.4				
NT2RF3000660	7.86	3.20	4.43	11.71	10.96	7.67	5.63	5.73	5.08				
NT2RF3000661	5.33	3.07	4.20	8.73	10.09	5.63	4.67	5.28	4.19				
NT2RF3000665	6.64	1.93	2.75	5.80	4.45	4.67	4.17	5.21	4.12				
NT2RF3000676	8.20	4.06	3.78	8.46	10.33	8.20	6.83	8.27	6.88				
NT2RF3000677	4.44	2.49	3.08	10.60	15.84	15.84	2.62	4.06	2.32	**	+		
NT2RF3000681	16.25	8.48	11.24	17.10	13.94	12.61	11.39	15.24	10.7				
NT2RF3000683	10.17	2.34	3.34	19.41	15.14	11.09	6.65	5.82	9.12	*	+		
NT2RF3000685	7.81	3.42	2.68	6.13	4.88	5.09	4.14	3.91	7.49				
NT2RF3000690	3.45	1.81	2.38	2.69	3.42	3.19	1.6	4.35	4				
NT2RF3000698	3.44	1.71	1.90	3.98	4.36	3.04	3.05	5.26	3.03				
NT2RF3000708	8.35	3.44	2.85	6.09	5.53	5.09	2.92	4.17	5.63				
NT2RF3000719	6.12	2.90	4.00	7.25	5.34	4.25	3.6	5.95	3.12				
NT2RF3000721	4.08	2.25	2.01	4.97	4.56	3.47	2.13	3.08	2.69				
NT2RF3000728	2.25	0.64	0.87	2.34	2.75	2.13	0.67	2.18	0.8				
NT2RF3000730	1.35	0.93	1.10	1.92	2.14	1.30	2.2	1.20	1.35				
NT2RF3000733	4.35	2.50	1.71	6.01	6.36	4.79	3.49	3.48	2.85	*	+		

99/180

【0740】

【表269】

1676

NTZRP000735	2.00	1.20	0.61	2.06	0.92	1.03	2.17	1.47	1.63										
NTZRP000736	3.46	3.21	3.33	4.48	4.58	3.34	3.43	2.28	2.96										
NTZRP000739	15.24	8.34	8.12	11.53	11.36	10.77	13.58	12.81	14.45										
NTZRP000742	15.14	9.63	9.98	14.05	14.60	13.15	13.09	11.17	13.06										
NTZRP000753	4.09	1.46	2.26	4.87	6.45	3.41	1.81	3.35	5.41										
NTZRP000759	4.36	3.02	3.28	9.27	10.72	9.10	9.4	9.92	12.65	**	+								
NTZRP000769	6.97	3.15	3.19	2.62	3.38	3.33	2.9	2.77	2.91										
NTZRP000815	3.08	1.87	2.78	5.08	5.91	5.79	4.34	3.06	3.31	**	+								
NTZRP000818	7.88	5.88	4.83	9.79	13.01	13.93	8.4	7.38	10.56	**	+								
NTZRP000829	6.70	4.35	2.57	15.50	20.24	18.97	5.35	5.01	5.38	**	+								
NTZRP000821	6.58	4.20	3.95	5.67	6.08	4.63	5.13	4.56	4.66										
NTZRP000825	0.66	0.26	0.38	1.28	1.09	2.20	0.44	1.29	0.44	*									
NTZRP000826	14.31	7.15	8.00	20.59	14.43	14.08	24	29.57	29.39	**	+								
NTZRP000836	8.67	4.78	5.47	15.61	15.21	9.41	7.61	8.33	8.85	*	+								
NTZRP000838	69.68	35.31	38.08	62.74	50.92	55.55	114.4	92.67	110.6	**	+								
NTZRP000859	3.11	1.70	2.32	2.00	3.56	1.87	3.03	1.90	2.5										
NTZRP000841	4.62	3.46	2.85	4.90	8.16	5.93	4.11	3.68	3.13										
NTZRP000845	4.22	3.31	3.16	4.56	7.12	4.56	4.69	3.53	11.01										
NTZRP000847	8.01	5.03	4.67	11.17	12.10	10.61	8.29	6.56	5.96	**	+								
NTZRP000848	4.58	2.34	3.27	5.39	6.00	5.09	3.72	3.05	4.52	**	+								
NTZRP000850	7.12	3.32	4.95	11.87	12.25	13.21	7.48	7.20	7.92	**	+								
NTZRP000852	2.41	2.02	2.14	2.50	3.10	2.98	1.15	2.04	2										
NTZRP000853	11.57	6.45	2.66	9.86	9.35	7.35	6.51	5.86	6.19										
NTZRP000861	12.29	5.70	6.74	20.57	26.68	20.53	8.96	8.46	14.99	**	+								
NTZRP000862	10.74	6.85	6.61	6.87	7.17	5.23	6.09	5.29	7.24										
NTZRP000865	2.61	2.77	1.86	4.46	4.70	3.49	3.05	2.82	3.22	*	+								
NTZRP000866	3.65	3.07	3.41	3.79	4.93	3.08	2.95	3.92	4.36										
NTZRP000868	6.63	4.07	4.55	4.52	6.19	4.40	5.59	4.36	6.01										
NTZRP000869	7.38	5.89	6.47	6.37	7.71	6.66	5.72	5.36	5.4										
NTZRP000871	2.80	1.69	2.21	3.13	2.44	2.63	2.19	2.91	2.3										
NTZRP000875	6.14	2.07	3.11	2.15	2.68	3.67	3.92	2.74	3.62										
NTZRP000885	3.27	2.20	2.57	3.83	6.39	6.13	3.73	2.67	3.88	*	+								
NTZRP000908	9.85	5.60	5.12	11.99	12.50	10.94	7.71	7.19	8.22	**	+								
NTZRP000901	5.01	2.45	2.11	6.45	8.36	6.11	4.49	5.69	7.42	**	+								
NTZRP000905	2.28	1.60	1.75	4.44	6.62	5.24	4.43	2.98	3.76	**	+								
NTZRP000904	2.30	1.61	2.05	2.19	1.89	3.67	2.54	3.22	2.14										
NTZRP000907	9.61	6.08	7.44	8.62	11.64	8.56	8.91	8.78	9.69										
NTZRP000913	7.70	2.80	3.71	8.25	8.06	6.91	5.87	6.50	4.94										
NTZRP000927	10.36	7.31	5.72	9.00	10.41	11.45	7.56	6.56	8.24										
NTZRP000919	5.76	4.04	3.02	5.13	7.71	4.25	4.75	4.65	6.91										
NTZRP000921	3.51	1.70	2.76	4.60	7.92	2.75	6.8	3.67	4.11										
NTZRP000942	9.61	5.52	5.34	12.62	14.38	12.46	6.8	6.33	7.24	*	+								
NTZRP000968	103.66	58.95	63.91	147.53	158.89	133.89	55.3	53.20	43.04	*	+								
NTZRP000974	3.04	1.59	2.65	3.97	5.03	4.21	2.71	3.66	2.41	*	+								
NTZRP000980	39.62	20.55	29.98	6.47	9.37	6.00	4.91	6.99	8.46	*	+								
NTZRP000984	5.29	4.18	5.73	10.16	10.11	7.87	6.25	8.85	4.44	**	+								
NTZRP000994	3.63	2.42	1.96	4.75	5.40	3.69	3.58	4.22	3.83										
NTZRP001001	3.47	2.25	3.10	3.83	2.41	2.13	2.68	3.98	2.58										
NTZRP001004	1.80	1.40	1.87	2.71	2.31	1.48	2.16	1.18	3										
NTZRP001007	4.63	2.03	2.66	14.00	6.75	8.49	6.39	6.25	5.07	*	+								
NTZRP001012	5.10	1.75	3.11	5.04	4.34	3.54	2.86	4.75	2.29										
NTZRP001042	5.71	3.43	4.72	5.27	4.96	3.88	3.98	3.86	2.98										
NTZRP001044	7.02	3.73	5.60	14.85	12.04	12.37	9.89	10.94	7.73	**	+								
NTZRP001048	2.35	1.96	3.94	3.25	4.98	4.26	3.16	2.56	3.24										
NTZRP001050	11.91	8.75	3.68	7.09	10.52	7.57	19.34	10.34	18.84										
NTZRP001055	19.61	12.87	10.53	9.87	9.64	7.47	11.2	7.71	10.89										

100/180

【表 270】

1677

1678

NT2RF3001057	8.67	4.03	5.93	19.26	14.18	12.30	8.42	6.94	7.2	*	+		
NT2RF3001061	5.88	4.01	4.14	7.73	9.70	8.03	5.42	6.19	4.88	*	+		
NT2RF3001069	9.78	4.93	5.43	13.99	17.62	14.76	9.74	9.96	12.86	**	+		
NT2RF3001074	8.31	4.57	4.04	11.86	10.34	7.95	6.59	7.36	7.45	*			
NT2RF3001078	5.34	2.26	4.49	9.51	7.77	7.53	5.94	3.60	3.02	*	+		
NT2RF3001081	3.83	2.45	4.20	6.12	3.89	6.40	3.56	5.22	3.4	*			
NT2RF3001084	5.54	2.82	2.70	2.36	4.10	1.78	2.85	2.45	3.36	*			
NT2RF3001095	1.93	1.69	1.44	3.80	3.49	3.25	2.25	2.83	2.47	**	+	*	+
NT2RF3001096	4.61	2.92	2.43	5.50	5.58	4.69	7.37	7.57	7.11	*		**	+
NT2RF3001097	9.61	7.40	9.00	12.56	12.16	11.92	6.67	6.88	9.12	**	+		
NT2RF3001107	6.04	4.02	3.50	4.89	5.87	4.23	3.8	4.49	5.02	*			
NT2RF3001109	6.26	3.05	4.30	3.18	4.47	2.65	2.28	2.72	1.85	*			
NT2RF3001111	4.22	3.38	2.92	4.15	5.15	4.60	4.36	4.69	3.98	*			
NT2RF3001112	28.16	25.89	21.28	24.06	17.68	24.85	10.06	10.73	13.22	*		**	
NT2RF3001113	1.79	0.99	0.62	1.34	2.23	1.54	1.11	1.24	1.25	*			
NT2RF3001115	3.88	1.85	2.25	7.26	3.45	2.57	3.4	3.91	4.67	*			
NT2RF3001116	3.94	1.69	1.56	4.63	3.42	2.74	3.29	4.13	4.33	*			
NT2RF3001119	9.02	6.38	5.74	6.52	9.40	7.53	6.04	4.43	6.5	*			
NT2RF3001120	11.82	5.87	8.94	18.20	12.33	18.08	8.42	9.14	10.96	*	+		
NT2RF3001124	3.38	2.35	3.59	5.64	8.45	7.51	8.01	7.65	6.3	*	+	**	+
NT2RF3001127	1.21	0.67	1.51	2.88	2.70	1.71	4.11	3.13	5.1	*	+	**	+
NT2RF3001133	7.23	4.12	5.49	7.95	8.82	7.67	4.57	6.00	4.72	*			
NT2RF3001140	2.84	1.04	1.66	3.30	3.99	3.19	1.56	2.10	3.38	*	+		
NT2RF3001147	7.62	3.19	3.51	4.05	4.82	4.29	0.77	2.87	1.63	*			
NT2RF3001150	5.19	1.79	3.13	6.49	3.73	3.77	3.8	3.66	4.52	*			
NT2RF3001152	2.12	0.44	0.89	1.69	1.74	1.98	1.83	2.36	2.08	*			
NT2RF3001155	6.90	4.51	4.25	3.69	4.69	3.75	1.73	3.87	3.96	*			
NT2RF3001156	2.47	1.68	1.60	2.59	3.59	3.31	2.51	4.84	4.65	*	+		
NT2RF3001159	12.19	5.40	5.34	9.00	9.95	7.35	6.84	6.11	6.14	*			
NT2RF3001170	7.10	4.60	5.72	9.66	13.09	10.69	5.5	6.89	3.71	*	+		
NT2RF3001176	9.51	3.49	2.75	17.93	12.62	10.20	6.88	5.97	13.3	*			
NT2RF3001195	6.18	2.83	2.96	6.39	10.42	3.54	4.18	5.32	5.17	*			
NT2RF3001209	29.33	14.29	10.79	23.50	28.08	21.04	16.75	19.48	15.61	*			
NT2RF3001214	6.63	3.46	3.32	9.82	10.42	9.38	3.48	5.63	3.56	**	+		
NT2RF3001216	4.48	3.19	3.11	7.11	8.39	8.87	2.58	5.22	3.57	**	+		
NT2RF3001221	1.19	0.31	0.47	1.55	1.56	1.10	1.01	2.22	0.86	*			
NT2RF3001226	7.00	2.58	2.80	4.50	5.21	4.34	3.95	5.75	3.9	*			
NT2RF3001230	2.86	1.59	1.71	4.14	3.19	2.63	1.59	3.61	2.59	*			
NT2RF3001232	4.81	1.38	0.57	1.61	2.09	1.97	2.63	1.53	0.99	*			
NT2RF3001236	1.71	1.43	0.80	2.59	2.82	2.72	3.58	2.05	2.31	**	+		
NT2RF3001239	2.21	1.46	1.67	2.79	2.29	1.43	3.36	2.12	2	*			
NT2RF3001240	2.39	2.60	2.79	4.11	6.20	4.44	7.84	6.72	4.74	*	+	*	+
NT2RF3001245	3.14	1.64	2.84	6.19	9.37	6.48	4.16	3.07	4.85	*	+		
NT2RF3001253	4.00	1.90	2.62	6.61	7.24	6.92	3.25	4.04	5.99	**	+		
NT2RF3001259	10.11	5.52	6.66	9.63	10.72	9.87	6.94	7.54	9.1	*			
NT2RF3001260	1.75	0.60	0.84	2.44	2.65	2.56	1.25	1.73	2.02	*	+		
NT2RF3001264	3.80	0.98	1.35	3.72	2.40	2.94	2.21	1.54	2.06	*			
NT2RF3001268	5.50	3.38	4.02	7.85	8.76	7.64	4.87	3.66	4.6	**	+		
NT2RF3001271	28.62	19.09	17.03	21.24	19.12	21.60	21.92	16.59	24.45	*			
NT2RF3001272	5.76	3.32	1.84	5.66	6.83	7.58	3.78	6.70	4.51	*			
NT2RF3001274	19.11	14.57	13.97	21.86	23.69	19.32	19.59	16.07	21.69	*	+		
NT2RF3001275	3.98	2.12	2.06	4.08	3.88	3.61	4.57	5.00	3.17	*			
NT2RF3001280	5.95	4.26	3.61	5.15	6.58	6.13	4	4.48	3.31	*			
NT2RF3001281	4.63	3.14	4.04	6.78	5.25	8.51	3.4	3.74	3.77	*	+		
NT2RF3001288	14.66	10.02	11.01	19.91	17.12	14.80	31.14	30.59	36.12	**	+		
NT2RF3001297	4.65	2.39	2.87	6.59	5.46	6.16	4.33	3.73	6.69	*	+		

101/180

【0742】

【表271】

1679

1680

NT2RF3001300	6.60	4.50	3.63	5.55	5.25	4.91	6.62	5.73	6.77										
NT2RF3001301	4.23	2.91	2.87	6.64	7.54	6.04	6.28	4.21	5.54	**	+								
NT2RF3001307	3.27	2.88	2.97	3.26	3.20	4.09	4.01	4.56	2.31										
NT2RF3001310	14.83	12.54	13.73	16.67	19.61	13.33	4.87	4.68	5.5			**	-						
NT2RF3001318	2.74	0.91	1.95	3.13	4.02	2.55	2.31	3.93	2.1										
NT2RF3001322	1.63	0.90	0.90	2.74	1.95	2.82	2.79	4.85	2.34	*	+								
NT2RF3001325	24.22	12.72	10.92	7.56	8.85	7.79	5.36	4.73	4.91										
NT2RF3001338	15.76	12.66	9.88	10.48	14.22	14.25	12.53	9.24	15.76										
NT2RF3001339	4.32	1.49	2.10	2.91	3.84	2.95	3.46	1.89	3.19										
NT2RF3001340	19.62	12.27	15.41	18.07	21.26	18.46	16.08	15.90	21.63										
NT2RF3001341	4.04	2.16	2.75	3.64	4.76	3.69	3.08	3.25	2.32										
NT2RF3001354	12.69	8.27	10.24	14.38	16.19	12.96	8.57	6.12	4.87										
NT2RF3001355	3.39	2.67	2.73	4.52	3.86	4.06	3.69	3.97	3.97	*	+	*	+						
NT2RF3001356	2.63	2.41	2.61	3.21	3.25	2.89	2.82	3.46	1.7	*	+								
NT2RF3001359	5.31	3.10	1.88	3.19	6.05	4.15	4.41	3.34	3.75										
NT2RF3001364	6.03	3.09	3.48	5.69	5.56	4.55	3.38	5.70	5.8										
NT2RF3001373	5.46	3.57	2.36	4.41	5.80	3.94	5.01	3.68	6.3										
NT2RF3001374	2.93	1.03	1.18	2.06	2.91	2.46	1.54	1.85	1.14										
NT2RF3001383	6.37	4.77	6.05	9.28	12.56	10.77	4.11	4.30	3.48	**	+	*	-						
NT2RF3001384	4.58	2.86	3.25	5.41	5.38	4.60	5.49	4.04	4.15										
NT2RF3001388	3.94	3.65	4.40	11.98	17.15	15.81	10.54	11.04	15.23	**	+	**	+						
NT2RF3001392	3.83	1.90	3.17	5.39	4.17	3.84	3.44	3.44	2.66										
NT2RF3001396	2.00	1.30	0.75	2.42	4.93	3.82	4.83	3.81	2.6	*	+	*	+						
NT2RF3001398	11.01	6.05	6.28	7.94	10.96	10.36	8.08	7.65	10.79										
NT2RF3001399	8.19	4.25	5.07	7.54	8.60	8.41	4.97	7.59	6.74										
NT2RF3001402	2.09	1.57	1.57	3.12	4.36	4.40	2.46	3.16	5.1	**	+								
NT2RF3001407	9.10	4.59	5.21	13.05	12.91	13.40	7.95	7.65	8.13	**	+								
NT2RF3001416	2.87	2.04	3.00	3.89	8.00	5.00	4.89	5.09	4.41			**	+						
NT2RF3001429	5.16	2.34	2.93	5.77	5.70	6.45	3.3	5.56	7.47	*	+								
NT2RF3001435	3.64	1.83	2.78	5.54	5.58	5.80	4.28	4.76	3.32	**	+								
NT2RF3001436	9.63	6.68	3.99	4.77	6.95	7.51	7.14	4.61	7.91										
NT2RF3001437	4.50	3.40	2.04	4.15	3.27	4.34	2.81	4.38	3.95										
NT2RF3001438	4.16	3.58	4.14	7.37	9.48	9.19	4.5	5.96	3.98	**	+								
NT2RF3001439	2.71	0.65	1.93	11.45	6.48	6.19	4.59	7.62	3.98	*	+	*	+						
NT2RF3001433	3.34	1.56	1.82	4.80	3.24	3.78	1.92	2.57	3.01										
NT2RF3001439	6.50	4.98	6.18	6.78	9.50	6.94	5.45	6.68	5.8										
NT2RF3001441	4.58	1.98	2.38	4.38	3.89	3.43	3.38	5.92	9.79										
NT2RF3001446	2.76	1.23	2.57	5.63	7.47	6.18	5.44	6.68	4.2	**	+	*	+						
NT2RF3001447	8.22	4.12	2.95	6.40	8.22	5.10	3.65	5.93	6.09										
NT2RF3001449	4.73	2.05	2.23	6.25	6.19	5.57	6.13	6.05	7.57	*	+	*	+						
NT2RF3001453	6.27	2.66	2.61	7.65	7.63	7.03	4.7	5.93	5.45	*	+								
NT2RF3001457	5.03	2.53	2.21	3.77	4.85	3.80	3.24	4.94	2.94										
NT2RF3001459	2.60	1.82	2.24	2.49	3.26	2.21	2.13	3.94	1.79										
NT2RF3001463	3.43	2.23	2.76	3.05	4.78	3.63	2.47	3.86	2.66										
NT2RF3001466	0.65	0.45	0.93	0.79	1.40	1.78	1.01	1.24	0.81										
NT2RF3001472	5.02	3.77	3.20	8.63	6.87	6.75	5.25	4.56	5.18	*	+								
NT2RF3001475	16.30	4.98	4.56	9.54	12.17	8.13	7.39	5.93	7.4										
NT2RF3001479	11.30	7.78	6.68	11.47	10.59	7.30	7.74	6.55	7.95										
NT2RF3001499	1.44	1.38	1.33	3.68	2.94	3.11	4.42	3.30	2.91	**	+	**	+						
NT2RF3001492	3.13	2.23	1.38	5.46	5.82	3.49	2.27	3.77	3.59	*	+								
NT2RF3001495	4.27	2.41	2.48	4.72	5.59	4.95	3.72	4.06	3.66	*	+								
NT2RF3001497	3.41	1.98	2.83	6.14	5.70	4.65	3.85	3.87	3.68	*	+								
NT2RF3001501	3.65	1.22	1.98	4.41	3.90	3.76	3.18	3.14	3.33										
NT2RF3001507	8.81	6.07	6.17	11.31	10.29	10.39	6.88	6.90	7.25	*	+								
NT2RF3001529	9.25	3.58	2.90	11.50	12.88	7.44	5.1	3.82	4.38										
NT2RF3001533	8.31	2.40	2.73	6.50	6.03	5.12	5.17	5.15	4.98										

102/180

【0743】

【表272】

1681

1682

NT2RF3001539	12.56	6.40	7.00	10.99	10.52	8.15	6.27	6.65	5.19										
NT2RF3001540	3.56	1.19	1.50	6.99	9.11	5.28	2.14	3.06	2.68	*	+								
NT2RF3001549	9.80	7.45	10.38	11.31	10.30	10.04	7.8	5.81	7.76										
NT2RF3001554	3.44	2.57	2.68	4.38	5.21	3.74	3.1	4.12	3.42	*	+								
NT2RF3001560	1.98	0.84	1.82	2.21	1.46	2.33	2.57	1.64	2.81										
NT2RF3001561	7.62	4.57	4.64	6.91	8.11	8.03	7.34	7.68	6.78										
NT2RF3001564	12.59	4.99	5.10	22.94	20.84	14.16	5.83	7.51	11.43	*	+								
NT2RF3001568	10.68	5.54	6.19	6.22	5.75	5.19	2.58	3.78	3.57										
NT2RF3001575	10.33	5.99	5.52	11.60	12.09	8.47	6.09	5.98	6.46										
NT2RF3001580	3.56	1.35	1.99	5.39	3.01	3.50	2.91	3.43	3.66										
NT2RF3001587	9.27	5.60	6.48	9.67	8.64	7.91	3.57	5.67	3.81										
NT2RF3001589	4.49	2.24	2.17	4.59	7.05	6.18	4.42	5.38	3.17	*	+								
NT2RF3001592	4.37	2.01	2.87	4.75	5.39	5.86	3.63	4.01	2.99	*	+								
NT2RF3001607	0.30	0.54	0.84	0.71	1.22	1.55	0.82	2.08	0.53										
NT2RF3001608	7.31	2.87	2.62	6.20	4.67	5.11	3.69	5.29	6.29										
NT2RF3001613	11.75	4.76	3.72	8.30	8.98	5.57	5.89	6.91	7.14										
NT2RF3001619	4.55	2.53	2.20	3.99	4.12	3.29	2.64	4.30	2.99										
NT2RF3001621	7.09	6.13	3.47	2.20	2.82	2.95	1.51	2.76	2.37										
NT2RF3001629	3.07	1.05	1.36	2.67	2.54	2.74	1.29	3.63	1.56										
NT2RF3001630	4.04	2.39	2.24	3.71	3.71	2.59	1.51	3.51	0.99										
NT2RF3001631	24.78	10.11	12.40	17.73	20.88	13.17	4.28	8.91	6.44										
NT2RF3001634	9.27	2.72	5.54	7.96	8.15	7.28	4.29	5.79	4.53										
NT2RF3001642	5.13	3.42	2.92	6.54	7.68	6.47	5.19	3.70	3.73	*	+								
NT2RF3001644	3.27	1.84	0.92	3.18	2.57	2.35	5	2.95	3.44										
NT2RF3001658	3.62	2.89	1.93	2.64	3.29	4.41	2.44	1.58	2.48										
NT2RF3001667	1.93	2.07	1.35	2.81	3.65	4.62	4.85	5.42	7.49	*	+	**	+						
NT2RF3001671	7.66	4.46	4.89	5.72	6.98	5.49	3.11	2.99	4.06										
NT2RF3001672	5.04	4.31	3.86	3.93	4.78	3.32	4.59	4.37	7.43										
NT2RF3001676	3.97	2.04	5.02	4.84	5.72	3.79	2.56	2.60	3.1										
NT2RF3001678	5.11	3.61	3.12	4.03	3.95	2.98	4.85	3.51	3.88										
NT2RF3001679	5.80	3.94	3.38	8.40	8.81	5.85	11	8.10	8.4	*	+	*	+						
NT2RF3001682	11.08	7.03	6.66	4.48	3.93	2.41	1.86	2.18	2.25	*	+	*	+						
NT2RF3001685	5.84	2.49	1.45	5.20	7.06	5.72	3.81	3.24	3.24										
NT2RF3001688	9.98	5.14	4.96	11.67	15.18	13.11	7.75	5.30	4.79	*	+								
NT2RF3001690	6.37	3.50	2.59	4.35	7.48	8.72	4.02	4.96	4.94										
NT2RF3001693	13.26	8.38	9.13	9.74	11.97	8.26	6.72	8.53	7.59										
NT2RF3001696	6.95	4.47	3.30	15.84	17.48	7.56	13.16	12.78	11.08			**	+						
NT2RF3001698	6.30	3.93	3.04	7.50	5.16	4.97	10.41	6.02	8.18										
NT2RF3001700	3.49	1.19	1.57	2.49	3.70	3.38	4.25	2.37	2.33										
NT2RF3001712	11.74	6.82	5.41	22.86	35.26	39.54	12.07	11.43	15.14	*	+								
NT2RF3001716	7.22	3.02	4.03	8.79	10.51	6.60	4.73	4.70	5.85										
NT2RF3001724	15.75	4.14	3.21	5.86	6.17	7.63	4.16	4.41	4.61										
NT2RF3001727	8.66	6.49	5.38	14.44	7.82	11.73	11.95	13.12	10.93			*	+						
NT2RF3001729	1.93	0.96	0.61	2.40	2.57	2.22	2.16	2.35	2.73	*	+	*	+						
NT2RF3001730	6.71	4.57	7.74	11.66	10.98	8.11	6.76	8.86	5.97										
NT2RF3001733	2.88	2.06	0.55	2.95	3.43	1.42	2.02	2.52	2.06										
NT2RF3001737	6.70	4.04	4.02	6.45	5.41	5.38	5.72	3.92	6.08										
NT2RF3001738	10.91	6.90	7.77	7.27	7.41	7.04	6.92	5.83	6.78										
NT2RF3001739	5.34	4.75	4.43	4.78	6.81	5.30	5.03	4.71	6.57										
NT2RF3001742	5.50	3.13	4.00	3.39	9.70	3.77	4.55	5.25	8.02										
NT2RF3001751	13.48	12.01	10.94	15.12	15.40	18.57	7.79	9.88	12.42	*	+								
NT2RF3001752	4.05	3.78	2.59	14.37	14.59	7.40	13.28	13.75	10.73	*	+	**	+						
NT2RF3001753	4.22	3.12	2.93	5.12	4.27	8.95	2.67	3.47	2.04										
NT2RF3001754	24.40	11.37	10.27	18.41	20.20	17.55	14.78	11.55	16.09										
NT2RF3001756	3.63	3.86	3.16	12.94	21.73	28.36	7.24	4.83	10.97	*	+								
NT2RF3001764	6.68	4.75	3.99	4.90	5.39	5.66	4.29	5.91											

103/180

【0744】

【表273】

1683

1684

NT2RP3001771	3.51	2.93	3.35	3.89	4.06	3.55	3.61	4.21	5.23										
NT2RP3001777	4.09	2.96	3.01	5.51	4.45	3.91	4.86	5.16	6										
NT2RP3001783	2.53	2.57	1.95	6.76	6.36	6.69	4.29	5.47	3.41	**	*	*	*	*	*	*	*	*	*
NT2RP3001792	5.75	4.70	5.90	6.11	8.15	9.14	6.11	4.96	5.99										
NT2RP3001799	4.41	4.21	3.75	7.39	9.01	7.29	5.88	7.01	5.73	**	*	**	*	*	*	*	*	*	*
NT2RP3001819	6.61	3.33	1.74	4.45	5.18	4.58	4.38	3.34	4.47										
NT2RP3001829	60.87	38.63	36.73	56.07	52.70	55.16	28.32	28.08	35.16										
NT2RP3001834	10.17	3.74	4.77	10.85	13.55	11.18	6.57	5.69	7.14										
NT2RP3001839	15.46	12.06	10.35	17.55	22.87	17.91	17.89	15.53	21.32	**	*	*	*	*	*	*	*	*	*
NT2RP3001844	3.92	4.22	4.08	8.68	8.00	8.70	4.83	4.18	5.54	**	*	*	*	*	*	*	*	*	*
NT2RP3001848	8.51	3.03	3.37	7.54	6.39	7.94	7.05	8.18	5.83										
NT2RP3001854	4.31	3.66	2.93	4.93	7.64	5.42	5.84	9.19	10.46										
NT2RP3001855	1.08	0.62	0.41	0.88	3.15	1.50	2.17	1.51	1.24										
NT2RP3001857	8.74	5.14	3.23	3.88	5.79	4.95	4.34	4.47	3.21										
NT2RP3001859	5.96	2.68	3.12	1.87	2.69	2.83	2.52	3.04	2.59										
NT2RP3001861	8.95	6.91	5.65	7.71	8.95	8.02	9.41	9.63	9.39										
NT2RP3001866	1.78	1.67	1.30	2.40	3.59	1.96	3.62	3.94	3.33										
NT2RP3001871	1.22	1.47	1.24	4.28	5.33	4.06	5.94	5.76	6.13	**	*	**	*	*	*	*	*	*	*
NT2RP3001874	2.39	1.48	1.04	1.60	1.73	1.49	2.15	3.07	2.44										
NT2RP3001878	1.89	1.50	2.48	4.52	7.04	3.00	1.74	2.47	2.05										
NT2RP3001883	4.23	3.76	3.61	4.08	6.00	8.45	4.94	5.08	4.08										
NT2RP3001896	3.95	2.31	1.26	3.38	7.80	4.28	4.49	2.83	4.64										
NT2RP3001898	12.61	5.06	3.64	6.11	6.18	5.92	8.68	7.13	11.31										
NT2RP3001899	5.05	3.28	2.34	3.69	5.19	3.08	2.74	3.58	3.91										
NT2RP3001901	12.98	8.89	8.12	8.50	8.51	10.47	8.45	6.54	7.26										
NT2RP3001915	6.53	3.55	4.50	3.73	7.04	4.19	2.46	3.27	3.28										
NT2RP3001924	0.32	0.45	0.32	1.03	1.16	1.31	0.6	2.68	0.45	**	*	*	*	*	*	*	*	*	*
NT2RP3001929	2.79	2.04	3.11	3.82	2.97	3.77	2.42	3.15	2.72										
NT2RP3001931	4.35	3.16	3.68	4.47	4.72	7.93	3.59	3.28	3.34										
NT2RP3001938	7.26	2.97	4.06	7.92	6.46	6.68	4	4.10	3.17										
NT2RP3001940	14.11	5.27	4.51	10.79	10.92	8.33	5.43	5.45	5.13										
NT2RP3001944	3.45	2.33	1.32	2.72	2.97	3.31	3.63	3.49	2.49										
NT2RP3001945	7.29	7.10	5.59	8.17	9.64	11.51	6.42	7.34	6.69	*									
NT2RP3001947	4.79	4.51	3.45	5.88	6.32	6.85	5.07	6.05	6.08	*	*	*	*	*	*	*	*	*	*
NT2RP3001949	2.69	1.52	2.67	4.00	3.55	3.46	2.68	2.84	2.52	*		*	*	*	*	*	*	*	*
NT2RP3001953	16.48	13.65	16.67	12.37	9.06	10.48	18.01	17.99	16.21	*		*	*	*	*	*	*	*	*
NT2RP3001954	5.28	2.86	2.85	5.44	4.55	3.42	3.76	3.67	4.11										
NT2RP3001956	24.22	13.29	14.18	28.43	28.08	22.94	14.79	12.62	14.22										
NT2RP3001967	7.52	2.65	2.30	9.80	9.24	5.06	8.63	5.51	4.88										
NT2RP3001969	7.99	4.86	4.65	5.70	7.31	4.72	3.47	2.46	4.31										
NT2RP3001976	7.58	3.73	3.57	8.43	12.72	10.69	5.69	4.81	4.65	*		*	*	*	*	*	*	*	*
NT2RP3001986	4.77	4.42	3.72	5.84	6.16	3.49	3.93	4.27	4.43										
NT2RP3001989	0.59	0.37	0.61	1.26	1.01	1.46	1.37	2.34	1.2			*	*	*	*	*	*	*	*
NT2RP3002002	4.38	2.14	1.97	6.96	7.70	8.52	3.16	5.19	5.28	**	*	*	*	*	*	*	*	*	*
NT2RP3002004	2.02	1.54	1.44	3.44	3.14	2.45	2.24	2.01	2.64	*		*	*	*	*	*	*	*	*
NT2RP3002007	2.30	1.16	1.11	2.63	2.31	2.50	1.77	1.64	1.85										
NT2RP3002014	4.46	3.07	2.32	5.12	6.41	4.59	6.11	3.83	4.25										
NT2RP3002015	7.60	4.06	4.17	6.58	5.55	3.85	6.25	4.00	5.51										
NT2RP3002033	1.85	1.50	1.64	2.80	2.86	2.23	1.62	2.56	2.12			*	*	*	*	*	*	*	*
NT2RP3002045	1.82	1.00	1.37	1.94	4.75	2.88	1.69	2.27	2.09										
NT2RP3002054	2.00	1.59	0.94	1.90	1.75	2.02	1.61	2.56	2.14										
NT2RP3002056	2.28	1.93	1.78	6.00	7.33	7.83	2.85	2.69	4.61	**	*	*	*	*	*	*	*	*	*
NT2RP3002057	1.99	1.12	1.41	3.14	2.25	1.70	1.48	2.86	1.82										
NT2RP3002061	16.71	9.57	7.36	24.61	19.84	16.31	10.52	9.51	8.46										
NT2RP3002063	2.33	1.47	0.86	3.09	3.51	2.69	1.64	2.02	3.16	*		*	*	*	*	*	*	*	*
NT2RP3002063	8.43	3.19	2.56	5.90	5.68	4.65	5.99	6.66	4.47										

1685

1686

NT2RF3002064	5.17	3.05	2.46	4.06	7.44	4.88	4.84	4.54	4.14				
NT2RF3002071	2.33	1.51	1.99	1.86	2.27	1.50	2.16	2.61	2.43				
NT2RF3002073	5.31	4.25	4.41	3.45	4.48	3.77	3.46	5.18	3.88				
NT2RF3002074	3.99	3.21	3.54	3.26	5.35	3.47	3.41	4.15	2.51				
NT2RF3002075	4.75	2.10	2.19	6.52	7.60	4.15	6.03	5.22	3				
NT2RF3002077	8.02	3.34	2.61	6.63	4.07	3.18	5.14	4.74	2.68				
NT2RF3002081	10.07	7.99	7.00	4.79	4.27	3.26	2.76	2.42	1.41	*	-	**	-
NT2RF3002086	4.94	3.90	3.43	7.01	9.40	7.91	6.79	5.61	5.45	**	+	*	+
NT2RF3002094	55.21	38.13	49.40	26.53	35.64	30.76	29.38	24.30	29.05	*	-	*	-
NT2RF3002096	2.03	2.45	2.09	2.34	2.63	1.70	2.31	1.94	2.22				
NT2RF3002097	4.81	2.56	2.66	7.07	9.45	4.39	4.28	5.92	5.09				
NT2RF3002098	1.30	1.49	2.04	3.02	3.52	2.73	1.86	1.80	1.76	*	+		
NT2RF3002102	4.48	2.97	2.73	5.04	5.32	5.08	5.06	4.28	4.93	*	+		
NT2RF3002106	5.41	2.39	2.38	9.26	7.89	8.90	6.1	3.83	3.57	**	+		
NT2RF3002108	6.53	3.49	4.50	3.88	5.75	3.58	3.09	4.07	3.18				
NT2RF3002109	11.23	5.02	4.28	16.19	18.27	13.88	14.35	12.01	12.31	*	+		
NT2RF3002110	23.37	14.84	16.48	34.91	29.71	40.33	23.01	21.75	23.48	*	+		
NT2RF3002113	11.63	9.01	7.67	6.51	7.35	7.47	7.32	7.10	6.45				
NT2RF3002120	1.55	1.48	1.08	2.91	3.24	1.92	2.33	3.13	2.18	*	+	*	+
NT2RF3002121	3.47	2.28	2.84	4.15	6.05	2.79	2.22	3.39	2.01				
NT2RF3002126	11.23	6.99	4.03	8.17	8.24	7.23	16.66	12.35	16.36			*	+
NT2RF3002128	13.16	6.63	6.22	10.39	10.13	7.09	9.73	7.03	10.29				
NT2RF3002130	7.94	5.84	4.52	8.35	9.12	8.25	8.69	6.14	9.87				
NT2RF3002133	7.90	4.13	2.94	10.10	13.02	11.57	10.36	9.95	10.86	**	+	**	+
NT2RF3002136	10.87	7.99	6.07	13.09	20.57	19.22	14.35	15.02	15.43	*	+	**	+
NT2RF3002140	4.41	4.46	5.24	5.99	5.61	7.54	7.49	4.80	5.22				
NT2RF3002142	7.81	6.29	3.94	14.63	15.34	11.73	11.3	15.25	13.24	**	+	*	+
NT2RF3002146	7.61	4.78	4.77	10.91	13.18	6.97	4.8	6.21	4.65				
NT2RF3002147	22.06	11.75	12.01	9.65	10.83	10.56	11.86	8.17	9.38				
NT2RF3002151	14.60	11.05	8.77	13.96	13.74	12.27	8.15	8.64	12.04				
NT2RF3002155	8.16	6.32	4.96	8.79	7.65	4.96	6.19	7.55	7.22				
NT2RF3002156	2.21	1.36	0.96	3.23	3.14	2.36	3.21	3.07	3.25	*	+	*	+
NT2RF3002160	3.98	3.19	1.94	3.32	4.52	5.20	4.3	1.89	4.12				
NT2RF3002163	18.81	11.61	12.16	18.87	21.42	15.74	12.51	9.05	10.05				
NT2RF3002165	6.12	5.16	5.75	6.38	8.10	3.82	6.23	5.63	7.23				
NT2RF3002166	5.72	3.53	1.35	2.95	5.16	3.30	2.3	3.24	3.17				
NT2RF3002173	5.34	3.03	2.78	9.80	6.20	7.21	5.06	5.00	4.94	*	+		
NT2RF3002174	5.68	2.49	1.67	7.29	8.21	9.12	9.02	7.21	12.43	*	+	*	+
NT2RF3002181	9.68	7.50	5.24	4.48	4.92	3.59	2.61	2.36	2.48			*	-
NT2RF3002185	3.81	2.37	1.77	2.88	7.87	3.22	3.57	3.44	2.54				
NT2RF3002193	7.51	6.09	4.76	5.28	9.69	7.23	6.2	5.26	7.9				
NT2RF3002204	2.89	2.47	0.95	9.64	8.53	14.75	4.05	4.67	4.6	*	+	*	+
NT2RF3002244	4.56	5.32	5.18	4.63	6.32	6.34	4.51	3.44	3.59			*	-
NT2RF3002248	8.18	5.72	5.54	14.10	16.32	12.91	11.02	10.26	11.54	**	+	**	+
NT2RF3002253	6.83	4.26	3.08	6.54	5.65	6.66	3.58	3.16	4.05				
NT2RF3002255	44.02	22.63	19.64	26.45	26.34	31.64	17.22	13.77	17.68				
NT2RF3002264	5.83	3.17	2.53	6.13	7.07	6.24	4.47	6.97	4.95				
NT2RF3002267	4.61	2.60	2.31	3.48	4.99	3.73	3.57	2.66	3.09				
NT2RF3002273	14.02	8.03	6.96	15.74	16.07	14.50	9.58	10.63	9.37				
NT2RF3002276	5.72	2.96	3.52	5.50	5.94	5.34	3.99	5.68	5.16				
NT2RF3002281	7.91	5.75	6.50	6.21	6.83	6.47	4.32	5.43	5.21				
NT2RF3002286	2.46	1.62	2.05	3.65	3.52	2.26	2.34	3.39	3.14				
NT2RF3002297	56.91	27.98	24.70	67.63	63.96	45.16	26.65	22.90	25.3				
NT2RF3002301	9.96	5.96	5.15	5.72	8.20	9.72	8.36	7.88	9.26				
NT2RF3002303	10.45	6.01	4.55	8.24	9.49	7.38	8.89	7.75	8.68				
NT2RF3002304	1.01	1.07	1.38	3.55	2.86	2.06	2.84	4.66	2.09	*	+		

105/180

【0746】

【表275】

1687

1688

NT2RF3002309	6.87	4.15	3.66	6.13	6.93	8.34	2.55	3.41	3.91										
NT2RF3002311	4.05	2.38	2.34	4.56	2.55	3.21	2.05	2.83	2.86										
NT2RF3002315	15.94	11.19	15.32	12.31	8.50	11.56	8.23	8.69	10.92			*	-						
NT2RF3002319	1.73	1.09	1.94	2.53	2.43	3.11	2.93	2.04	2.66	*	+								
NT2RF3002324	9.27	3.66	3.72	5.93	9.44	5.66	4.2	5.07	4.43										
NT2RF3002330	9.95	5.32	3.76	4.42	7.75	7.05	6.63	6.18	5.42										
NT2RF3002333	17.93	13.63	12.33	10.81	13.83	11.53	26.44	20.51	21.61			*	+						
NT2RF3002337	2.63	1.45	1.52	1.90	1.94	2.01	1.38	3.21	2.65										
NT2RF3002343	15.59	10.64	11.07	10.92	13.50	7.96	9.5	11.72	10.96										
NT2RF3002343	4.86	3.15	3.42	8.66	7.27	7.64	5.82	6.21	6.54	**	+	*	+						
NT2RF3002351	2.14	1.87	1.48	1.52	1.49	1.39	1.37	2.50	1.29										
NT2RF3002352	3.51	2.49	2.09	6.56	3.41	4.41	3.67	4.42	2.26										
NT2RF3002353	8.54	2.87	2.50	5.68	7.92	6.04	5.65	4.24	3.09										
NT2RF3002362	10.04	4.71	5.05	6.95	8.81	7.91	8.38	7.04	7.67										
NT2RF3002363	5.45	3.22	2.99	4.20	6.31	4.65	3.29	3.42	4.78										
NT2RF3002377	6.53	3.54	3.81	6.50	6.48	4.79	3.11	4.43	2.57										
NT2RF3002377	16.05	6.92	7.02	15.78	13.73	11.15	9.35	6.37	9.19										
NT2RF3002384	3.83	2.35	2.55	5.43	6.35	4.75	5.11	5.17	5.17	*	+	**	+						
NT2RF3002397	1.88	2.06	1.00	2.28	2.42	2.35	2.43	3.20	2.26										
NT2RF3002399	38.89	13.57	16.73	24.89	24.11	20.07	10.95	10.34	11.58										
NT2RF3002402	14.13	6.06	6.64	3.90	7.46	3.60	5.13	2.47	3.86										
NT2RF3002404	2.69	1.41	1.51	4.63	5.57	6.95	5.03	5.62	5.49	**	+	**	+						
NT2RF3002410	16.74	9.36	8.24	14.55	17.40	14.68	7.71	8.16	9.6										
NT2RF3002411	5.72	3.09	2.66	5.44	3.76	4.39	3.87	3.60	4.64										
NT2RF3002414	15.70	13.46	15.51	17.50	19.84	20.94	20.31	15.95	17.64	*	+								
NT2RF3002430	5.62	3.03	3.26	4.15	6.68	5.69	3.6	5.22	5.76										
NT2RF3002440	3.21	1.91	1.95	4.68	4.12	2.16	3.43	3.57	3.52										
NT2RF3002454	5.75	3.63	2.88	8.65	10.72	8.12	4.17	6.41	5.11	*	+								
NT2RF3002455	5.96	2.60	2.61	5.44	7.86	5.02	4.61	3.98	4.33										
NT2RF3002456	19.55	5.82	6.70	24.00	22.06	18.49	6.98	7.59	13.81										
NT2RF3002463	10.35	5.72	4.60	11.65	13.73	9.93	5.45	7.13	8.04										
NT2RF3002469	4.02	2.04	2.37	7.68	7.85	6.75	5.57	6.12	6.98	**	+	**	+						
NT2RF3002470	34.16	21.24	23.62	26.50	31.46	31.78	25.6	23.51	18.11										
NT2RF3002484	4.96	4.07	3.20	7.26	8.04	8.64	6.14	7.06	7.03	**	+	*	+						
NT2RF3002491	2.02	0.31	0.77	1.88	1.82	1.66	1.70	2.17	2.19										
NT2RF3002494	5.69	5.46	5.09	5.37	5.09	4.28	11.1	14.53	16.58			**	+						
NT2RF3002497	7.34	2.87	2.34	7.23	5.25	4.45	4.45	4.17	5.52										
NT2RF3002500	6.11	2.15	1.67	4.34	5.06	2.16	2.18	2.29	5.42										
NT2RF3002501	11.25	5.11	3.44	6.23	6.00	5.47	2.88	5.58	5.46										
NT2RF3002512	7.00	3.28	2.78	5.82	6.08	6.36	2.87	4.61	8.18										
NT2RF3002529	3.20	3.16	1.84	7.16	9.33	8.45	4.14	4.40	5.49	**	+	*	+						
NT2RF3002533	7.52	4.47	4.21	12.54	12.31	10.84	8.33	13.60	12.28	**	+	*	+						
NT2RF3002539	6.08	4.61	2.98	8.67	11.27	7.39	2.77	5.22	3.99	*	+	*	+						
NT2RF3002540	2.20	1.79	1.19	3.09	3.15	2.53	2.67	2.99	2.88	*	+	*	+						
NT2RF3002543	14.24	6.52	5.35	11.36	10.19	12.56	10.27	10.96	8.43										
NT2RF3002545	4.03	2.04	1.37	6.55	5.22	5.90	5.61	3.59	2.71	*	+								
NT2RF3002549	2.56	1.25	0.83	5.75	4.78	6.90	5.63	3.95	4.81	**	+	*	+						
NT2RF3002553	2.93	2.06	2.41	3.32	5.85	3.49	4.06	3.68	4.2			**	+						
NT2RF3002558	7.05	4.19	4.48	9.57	11.91	11.02	10.69	7.40	9.14	**	+	*	+						
NT2RF3002563	4.40	2.70	2.23	5.52	4.89	4.10	2.94	2.79	3.23										
NT2RF3002566	4.15	3.12	3.18	4.65	4.50	3.46	4.21	2.25	2.13										
NT2RF3002571	1.43	0.64	1.11	2.38	2.79	1.21	2	1.13	1.01										
NT2RF3002572	5.68	2.77	2.24	4.20	4.73	4.39	4.25	2.73	1.87										
NT2RF3002573	12.53	5.63	5.03	10.21	9.69	14.18	6.47	7.06	5.93										
NT2RF3002577	16.44	10.30	7.27	9.02	16.56	19.10	11.76	11.75	13.42										
NT2RF3002579	5.14	1.77	2.75	2.43	7.06	4.13	4.98	5.48	3.32										

106/180

【0747】

【表276】

1689

1690

NT2RF3002582	12.31	7.23	7.62	9.16	12.52	12.19	7.07	6.53	8.27				
NT2RF3002587	2.59	1.37	0.54	2.46	2.67	3.02	1.24	1.89	1.22				
NT2RF3002590	10.29	5.66	7.55	5.34	4.92	3.70	2.44	4.90	2.27		*	-	
NT2RF3002602	2.82	1.08	1.45	3.79	2.37	2.51	2.16	2.20	1.92				
NT2RF3002603	23.80	12.85	10.63	16.77	16.77	18.88	33.04	20.98	28.78				
NT2RF3002621	5.83	2.17	2.11	2.73	3.73	3.84	3.77	3.43	4.67				
NT2RF3002623	6.46	4.71	3.37	7.18	6.32	5.80	5.41	4.46	6.55				
NT2RF3002624	1.38	1.46	0.86	2.16	2.27	1.71	1.92	2.31	2.23	*	*	*	+
NT2RF3002628	3.88	4.12	4.54	3.93	5.98	4.39	6.01	5.25	6.35		*	*	+
NT2RF3002639	17.56	11.86	13.81	23.77	21.74	24.60	15.1	15.62	16.2	**	+		
NT2RF3002631	0.65	0.54	0.71	0.74	2.00	0.23	0.47	2.10	1.77				
NT2RF3002647	6.35	4.67	4.32	5.81	4.61	3.54	2.45	3.29	2.94		*	-	
NT2RF3002649	13.39	5.95	5.65	10.41	9.34	8.49	5.95	5.93	9.13				
NT2RF3002650	6.81	4.69	4.82	5.81	7.89	6.12	6.83	5.78	9.56				
NT2RF3002652	5.20	4.74	1.12	4.44	5.82	4.44	3.42	3.65	3.38				
NT2RF3002654	16.99	10.82	13.04	8.59	8.02	5.74	6.46	6.13	9.06	*	-	*	-
NT2RF3002657	6.11	3.63	4.64	10.15	11.45	6.16	9.57	10.27	10.97		**	+	
NT2RF3002659	1.43	1.66	1.88	2.50	3.07	1.94	1.45	2.43	1.88				
NT2RF3002660	6.69	4.61	2.72	7.71	9.95	6.32	4.86	5.91	5.04				
NT2RF3002663	2.95	2.45	2.08	3.55	3.38	2.69	2.33	2.32	1.43				
NT2RF3002664	4.14	2.04	1.66	3.83	4.46	3.08	3.81	2.61	3.84				
NT2RF3002667	10.84	11.80	12.31	7.37	13.24	10.35	2.54	3.53	3.86		**	-	
NT2RF3002671	4.10	3.38	2.05	3.68	4.13	3.09	3.64	4.14	3.95				
NT2RF3002683	6.85	6.11	3.50	9.41	10.82	9.25	7.6	6.54	14.33	*	+		
NT2RF3002684	2.31	2.12	2.06	2.65	2.46	1.95	3.43	3.91	2.52				
NT2RF3002687	0.81	0.83	0.64	1.63	2.27	2.37	2.18	2.59	1.3	**	+	*	+
NT2RF3002688	1.90	1.35	1.30	2.68	10.84	4.31	2.62	3.98	4.96		*	+	
NT2RF3002698	1.70	1.54	2.28	2.37	1.97	1.69	2.37	4.37	2.27				
NT2RF3002701	9.13	4.28	3.80	7.31	8.31	6.47	5.76	5.84	9.76				
NT2RF3002705	21.78	18.18	17.66	50.09	57.33	55.80	17.31	19.57	25.8	**	+		
NT2RF3002708	8.43	3.13	4.23	10.00	12.33	16.86	6.66	9.06	8.15	*	+		
NT2RF3002711	10.69	7.85	6.27	14.28	17.41	10.11	7.22	6.34	9.71				
NT2RF3002712	75.48	54.09	63.05	72.21	59.25	49.90	55.73	52.68	50.32				
NT2RF3002713	1.12	1.39	0.99	1.79	1.94	1.51	1.51	1.64	2.24	*	+		
NT2RF3002711	4.73	3.29	3.45	5.55	8.69	5.41	5.47	5.66	7.4		*	+	
NT2RF3002723	18.60	15.91	19.67	21.10	20.78	20.71	21.26	14.74	13.19				
NT2RF3002723	20.89	13.71	12.73	18.63	26.94	25.35	23.58	19.98	24.35				
NT2RF3002737	10.83	5.85	5.46	7.36	8.93	8.81	7.12	8.21	8.27				
NT2RF3002738	3.06	2.31	2.46	3.88	2.93	4.58	4.14	4.86	3.57		*	+	
NT2RF3002742	78.11	50.55	39.19	56.71	49.99	44.98	24.65	24.79	19.15		*	-	
NT2RF3002744	1.91	1.57	1.49	3.37	4.81	3.15	4.58	3.73	2.77	*	+	*	+
NT2RF3002756	2.31	1.24	1.63	1.83	2.14	1.21	1.7	1.60	2.11				
NT2RF3002757	4.69	3.13	4.35	7.14	8.49	8.18	8.15	8.37	8.37	**	+	**	+
NT2RF3002758	7.65	5.42	7.31	13.02	12.93	12.57	12.33	13.43	11.46	**	+	**	+
NT2RF3002762	17.62	11.52	8.02	10.66	16.28	10.88	8.09	6.08	11.34				
NT2RF3002763	5.98	3.76	3.67	4.32	6.42	5.16	4.76	6.11	4.92				
NT2RF3002770	6.69	2.71	1.54	4.12	4.84	3.63	3.88	6.30	6.14				
NT2RF3002771	4.19	4.34	2.59	8.14	7.86	8.58	10.72	12.24	8.84	**	+	**	+
NT2RF3002785	3.87	2.70	2.07	1.69	2.61	1.77	0.79	2.12	2.01				
NT2RF3002790	2.54	1.59	2.82	4.68	4.85	6.90	3.49	4.63	2.59	*	+		
NT2RF3002792	2.06	0.55	1.55	2.25	2.19	2.80	1.65	2.16	2.21				
NT2RF3002801	3.39	2.62	3.03	5.62	4.43	4.91	3.26	3.08	2.61	**	+		
NT2RF3002802	9.76	4.91	4.56	5.83	7.90	5.66	5.83	5.98	7.36				
NT2RF3002810	2.05	2.04	1.36	1.85	2.29	2.16	2.36	3.68	3.36		*	+	
NT2RF3002818	1.54	1.82	1.16	0.90	1.59	1.73	1.13	2.06	1.73				
NT2RF3002821	17.00	12.39	12.28	12.51	17.54	13.78	7.96	8.86	8.91		*	-	

107/180

[0748]

[表277]

1691

1692

NT2RF3002323	1.32	1.08	1.04	1.83	2.17	1.81	1.57	3.57	2.5	**	+		
NT2RF3002324	7.13	4.05	4.87	6.63	6.04	8.47	4.09	5.57	4.15				
NT2RF3002329	3.03	2.45	2.63	5.74	5.50	4.90	3	3.82	3.79	**	+		
NT2RF3002331	3.87	3.21	2.77	3.69	2.99	3.89	2.66	2.74	2.29				
NT2RF3002336	14.03	6.74	6.74	9.92	15.02	8.10	13.6	10.55	13.13				
NT2RF3002345	6.06	2.27	2.32	3.35	4.67	5.99	2.22	2.92	5.24				
NT2RF3002352	2.14	1.57	1.15	1.52	1.72	1.72	1.78	2.42	2.44				
NT2RF3002361	4.05	2.12	1.50	1.55	2.01	4.44	1.39	3.44	3.12				
NT2RF3002369	6.92	5.64	4.79	4.48	4.94	3.03	3.48	5.21	5.99				
NT2RF3002374	3.62	2.41	3.09	2.41	2.83	2.25	3.7	5.14	4.58				
NT2RF3002376	6.38	5.46	5.19	8.34	12.34	10.89	6.16	7.19	7.18	*	+		
NT2RF3002377	4.36	2.55	2.24	6.28	5.72	7.39	4.17	3.78	4.69	*	+		
NT2RF3002387	2.31	2.06	1.28	2.41	6.33	3.71	2.23	1.91	2.99				
NT2RF3002390	4.63	3.12	1.94	6.79	7.22	4.89	6.77	4.56	5.42	*	+		
NT2RF3002392	13.48	7.11	7.49	17.13	16.57	10.16	8.66	6.18	6.66				
NT2RF3002399	33.33	17.88	18.92	24.91	27.67	27.33	23.19	23.81	25.55				
NT2RF30023911	2.05	1.51	2.25	2.06	2.34	3.42	1.9	2.88	2.46				
NT2RF30023948	2.87	2.05	2.73	3.15	3.80	3.22	3.02	3.24	4.14				
NT2RF3002393	2.95	2.20	2.80	3.91	2.99	2.13	3.94	4.99	3.35				
NT2RF30023955	3.21	2.28	2.19	2.68	3.66	2.17	2.8	4.04	3.2				
NT2RF3002398	5.15	1.89	1.75	8.63	9.49	5.11	5.86	5.70	7.9				
NT2RF3002399	8.37	4.79	4.07	7.09	7.89	5.99	3.82	5.59	8.02				
NT2RF3002372	2.45	1.77	1.17	3.30	4.53	6.41	2.37	3.50	4.2	*	+		
NT2RF3002378	3.51	1.12	0.76	1.57	2.29	1.16	1.76	2.22	2.49				
NT2RF3002383	2.09	1.72	1.47	2.93	4.10	4.53	1.5	4.04	1.42	*	+		
NT2RF3002385	2.93	1.24	0.64	1.80	1.57	1.56	1.03	3.24	1.64				
NT2RF3002388	3.04	1.50	1.33	2.69	2.87	2.12	2.09	2.69	1.72				
NT2RF3003090	5.52	4.04	3.47	8.75	7.05	6.47	5.37	5.35	7.11	*	+		
NT2RF3003098	3.30	1.49	1.41	3.13	2.40	2.15	3.61	1.58	2.05				
NT2RF3003012	5.75	2.52	2.34	2.71	2.38	1.98	3.89	1.73	1.65				
NT2RF3003015	3.67	2.39	1.41	2.11	1.98	2.12	2.64	2.73	1.76				
NT2RF3003018	5.19	3.49	2.94	3.09	5.88	7.34	2.45	3.41	8.68				
NT2RF3003023	4.42	2.89	2.76	3.64	5.83	5.34	3.92	2.05	3.21				
NT2RF3003029	5.92	3.71	3.59	6.44	6.11	4.11	7.41	7.78	5.42				
NT2RF3003033	8.58	6.19	7.17	18.73	18.81	11.60	10.2	11.99	14.12	*	+	*	+
NT2RF3003041	0.23	0.21	0.07	0.41	0.42	0.07	0.35	0.34	-0.17				
NT2RF3003044	7.25	3.53	3.53	7.47	6.31	4.80	5.47	4.15	4.63				
NT2RF3003047	14.58	8.48	8.68	11.39	12.06	11.40	11.77	9.28	11.88				
NT2RF3003050	6.53	2.71	3.77	5.22	5.47	3.84	3.66	4.93	4.39				
NT2RF3003053	17.07	9.71	8.94	14.88	15.92	20.90	14.19	12.88	11.32				
NT2RF3003059	2.32	1.74	2.11	2.95	2.30	1.48	1.32	1.45	1.42		*		
NT2RF3003061	4.13	2.99	2.62	3.51	4.22	2.44	3.64	4.14	3.12				
NT2RF3003068	7.07	5.01	4.05	8.08	8.01	6.86	3.94	4.27	5.35				
NT2RF3003071	7.18	5.69	5.64	19.53	14.10	9.02	3.53	6.99	4.86				
NT2RF3003076	20.24	13.69	11.73	17.25	17.10	20.23	12.75	12.44	19.06				
NT2RF3003078	6.31	1.99	2.60	4.81	6.42	5.61	4.7	3.16	4.19				
NT2RF3003081	5.58	3.59	4.40	7.90	10.09	9.19	6.17	5.95	6.74	**	+	*	+
NT2RF3003089	4.22	2.78	2.81	6.19	7.29	6.41	3.45	3.07	3.62	**	+		
NT2RF3003097	2.80	1.80	2.13	3.12	4.85	3.19	4.18	3.63	2.96		*	+	
NT2RF3003098	3.43	1.98	2.02	2.28	3.12	2.15	2.19	2.67	2.43				
NT2RF3003101	5.48	5.07	5.35	6.08	7.76	5.95	4.99	7.09	5.03				
NT2RF3003109	14.31	7.48	6.90	18.37	16.28	14.44	15.14	17.70	14.3				
NT2RF3003121	150.76	6.07	4.19	6.53	16.16	5.37	32.59	5.96	252				
NT2RF3003133	6.04	4.14	3.20	8.62	13.38	13.91	5.16	4.56	8.91	*	+		
NT2RF3003137	10.77	5.97	6.19	4.43	6.11	4.14	3.41	3.49	4.4				
NT2RF3003138	5.81	4.35	3.40	6.66	5.96	5.72	1.99	2.76	2.93				

108/180

[0749]

[表278]

1693

1694

NT2RF3003139	2.43	1.97	1.82	4.72	6.45	3.81	3.26	3.26	4.15	*	+	*	+
NT2RF3003145	2.66	3.16	2.32	3.58	4.86	4.52	5.45	3.67	3.72	*	+	*	+
NT2RF3003150	4.45	3.91	3.35	3.70	3.28	5.66	5.36	4.59	2.96				
NT2RF3003157	15.45	8.45	11.15	23.44	27.58	18.86	11.74	13.90	10.21	*	+		
NT2RF3003185	3.41	2.15	1.16	2.42	3.21	3.33	3.63	2.51	4.07				
NT2RF3003190	5.13	4.24	4.83	11.32	20.02	13.42	6.1	6.95	8.42	*	+	*	+
NT2RF3003197	3.94	1.73	2.04	2.63	7.18	5.13	2.76	3.02	4.74				
NT2RF3003203	10.74	6.48	7.57	9.78	9.35	10.34	12.74	12.49	16.29			*	+
NT2RF3003204	5.10	4.07	4.28	9.44	9.51	9.35	6.59	6.58	5.8	**	+	*	+
NT2RF3003210	2.87	2.26	2.76	4.58	4.94	5.68	4.02	4.31	4.86	**	+	**	+
NT2RF3003213	3.99	3.41	3.08	11.16	9.44	5.92	5.65	5.21	4.76	*	+	**	+
NT2RF3003213	3.64	1.51	1.06	6.12	6.44	4.09	4.51	3.54	3.74	*	+		
NT2RF3003224	4.97	2.24	2.03	3.13	4.35	3.48	1.88	2.89	5.66				
NT2RF3003226	6.57	4.20	3.82	3.03	7.40	7.29	3.35	4.03	3.53				
NT2RF3003230	5.88	2.80	3.00	5.34	6.53	3.95	6.24	6.52	4.98				
NT2RF3003235	5.68	3.50	3.55	11.57	10.92	8.51	10.86	10.22	8.85	**	+	**	+
NT2RF3003242	2.60	1.56	1.56	2.17	2.65	0.82	2.88	3.62	2.34				
NT2RF3003251	6.96	4.06	5.58	8.26	9.86	10.16	5.03	5.10	5.01	*	+		
NT2RF3003253	3.92	3.17	2.70	4.36	6.32	3.73	3.3	3.53	3.19				
NT2RF3003258	4.44	4.88	3.51	5.73	7.67	6.20	6.76	5.52	8.07				
NT2RF3003260	10.73	5.21	4.49	3.79	7.69	5.80	4.33	3.45	7.99				
NT2RF3003264	3.02	3.32	2.19	15.38	18.88	12.82	6.5	5.90	7.82	**	+	**	+
NT2RF3003273	3.18	1.91	3.15	2.64	2.58	3.24	1.86	3.56	1.93				
NT2RF3003278	3.16	1.06	0.85	1.38	1.88	2.32	0.32	2.57	2.1				
NT2RF3003280	11.26	9.07	8.30	12.96	14.31	12.01	8.63	10.92	9.88	*	+		
NT2RF3003283	2.12	1.63	1.57	3.75	3.52	2.64	2.53	3.71	3.58	*	+	*	+
NT2RF3003280	6.74	3.39	5.29	8.39	9.77	12.47	5.55	7.58	4.52	*	+		
NT2RF3003301	3.39	1.66	2.31	5.80	5.15	3.88	3.51	3.63	2.51	*	+		
NT2RF3003302	4.39	1.94	0.70	3.91	4.34	3.52	1.87	2.40	2.1				
NT2RF3003311	6.06	3.51	2.81	1.70	1.60	1.58	1.38	2.35	2.23				
NT2RF3003312	2.65	1.71	1.08	1.61	2.31	2.14	2.34	3.94	2.4				
NT2RF3003313	2.10	1.55	1.28	2.78	3.32	3.29	2.46	3.52	2.12	**	+		
NT2RF3003327	4.75	3.06	2.77	5.48	4.57	3.91	2.76	4.36	2.87				
NT2RF3003330	2.85	1.28	1.93	2.62	3.38	1.73	2.22	3.78	2.76				
NT2RF3003344	2.79	2.00	1.76	2.66	2.98	3.04	2.26	1.95	1.8				
NT2RF3003346	5.06	3.51	3.24	6.69	7.03	5.74	4.23	5.12	4.21	*	+		
NT2RF3003349	9.03	3.41	4.20	7.42	11.99	8.27	4.03	4.39	5.81				
NT2RF3003353	2.34	1.65	0.86	3.37	3.35	2.15	1.51	2.11	2.73				
NT2RF3003354	28.51	16.58	19.06	32.92	34.54	31.72	24.06	25.56	26.43	*	+		
NT2RF3003368	4.73	3.35	3.40	3.00	5.12	6.89	5.78	4.93	4.85				
NT2RF3003375	7.10	4.96	7.12	8.55	8.55	5.98	2.32	4.29	4.97				
NT2RF3003377	7.20	4.93	4.97	2.66	4.68	3.75	3.7	3.85	3.56				
NT2RF3003384	2.46	2.07	1.01	3.30	3.65	2.66	3.02	2.86	2.88				
NT2RF3003389	5.42	4.79	5.32	4.48	4.42	6.30	6.9	4.85	5.01				
NT2RF3003396	9.36	4.73	3.86	5.45	9.23	6.23	6.71	5.33	6.72				
NT2RF3003403	3.05	1.65	1.51	5.41	4.67	5.69	2.27	2.49	2.78	**	+		
NT2RF3003409	2.84	1.35	2.12	3.28	3.13	1.88	2.79	2.04	3.14				
NT2RF3003411	8.55	4.92	6.03	7.49	10.91	12.20	6.24	6.43	9.99				
NT2RF3003428	4.15	2.44	2.36	6.31	7.10	8.42	3.61	4.27	6.2	**	+		
NT2RF3003425	3.63	2.52	1.95	2.10	3.46	2.62	1.83	3.37	4.24				
NT2RF3003426	9.31	6.11	5.45	12.48	10.53	11.34	9.4	9.12	8.46	*	+		
NT2RF3003427	8.99	4.74	5.99	5.37	7.25	6.52	7.72	5.96	8.43				
NT2RF3003433	9.63	4.28	3.87	11.80	10.94	8.04	5.78	4.80	7.07				
NT2RF3003437	18.34	7.27	6.59	15.83	19.21	15.84	12.02	13.41	15.15				
NT2RF3003448	6.95	4.49	2.68	9.13	8.95	5.55	5.56	5.02	6.11				
NT2RF3003455	8.08	3.54	2.51	10.14	9.73	10.02	5.84	8.37	5.46	*	+		

109/180

【0750】

【表279】

1696

NTZRP3003462	4.12	2.91	3.40	4.80	6.31	3.84	3.87	5.08	5.3								
NTZRP3003464	2.09	1.93	2.23	2.69	2.89	1.33	2.08	3.11	2.15								
NTZRP3003466	3.14	2.14	3.25	2.94	3.46	3.12	4.23	4.89	4.48								
NTZRP3003473	89.05	73.91	85.12	31.82	60.74	51.48	52.07	37.37	37.95	*	-	**	-				
NTZRP3003474	3.72	1.64	1.41	2.81	4.68	2.60	1.76	1.83	5								
NTZRP3003475	5.61	2.84	3.02	4.26	5.48	3.96	3.12	2.86	5.38								
NTZRP3003490	2.57	1.77	0.80	2.92	3.66	2.60	1.94	2.99	7.23								
NTZRP3003491	3.83	1.51	1.56	3.52	3.19	3.71	1.08	3.23	2.23								
NTZRP3003493	32.32	24.24	22.86	18.58	22.23	21.78	10.29	22.93	16.25								
NTZRP3003508	1.40	1.72	1.09	3.53	3.58	2.03	2.23	3.61	2.95	*	+	*					
NTZRP3003527	2.93	1.02	3.29	2.26	3.40	1.33	1.5	4.37	4.13								
NTZRP3003532	6.83	4.04	4.22	15.20	17.07	14.08	8.23	6.65	7.28	**	+						
NTZRP3003536	1.58	1.03	3.00	1.85	1.07	0.98	1.62	1.27	0.97								
NTZRP3003538	2.90	2.77	1.64	5.15	3.92	4.74	3.97	3.71	2.81	*	+						
NTZRP3003543	4.72	4.39	3.25	5.41	8.08	7.02	5.2	5.69	1.98	*	+						
NTZRP3003549	2.71	2.81	2.37	2.41	3.29	4.08	3.3	2.30	1.66								
NTZRP3003552	1.05	1.06	0.00	1.19	1.29	1.21	0.42	0.50	0.79								
NTZRP3003555	7.69	3.49	4.38	7.36	8.58	9.29	5.4	5.33	4.02								
NTZRP3003559	2.48	1.02	1.13	2.42	2.15	3.37	1.46	1.90	0.56								
NTZRP3003564	6.10	3.28	3.23	6.06	5.72	4.12	4.46	3.78	4.48								
NTZRP3003572	4.33	3.51	2.66	3.48	5.50	3.26	3.67	4.32	2.39								
NTZRP3003576	14.59	6.63	6.37	16.23	21.96	19.84	10.82	10.96	8.97	*	+						
NTZRP3003587	15.06	8.22	7.98	8.40	8.95	10.51	3.86	6.51	3.33								
NTZRP3003589	14.90	11.19	8.98	10.98	18.16	17.00	16.77	16.70	14.61								
NTZRP3003592	6.07	3.40	4.66	3.72	5.45	5.40	3.54	4.53	3.89								
NTZRP3003593	5.28	1.73	2.13	3.76	3.86	7.00	3.06	4.00	2.78								
NTZRP3003614	14.05	8.27	10.10	10.29	8.15	9.17	8.06	7.21	4.02								
NTZRP3003621	3.29	1.07	1.69	2.23	2.27	2.45	2.08	2.77	2.99								
NTZRP3003625	11.53	5.52	5.48	9.50	9.71	7.13	7.18	5.14	5.56								
NTZRP3003637	12.05	7.44	6.80	53.97	42.81	41.76	14.96	15.18	18.14	**	+	*	+				
NTZRP3003636	5.65	3.72	2.95	5.93	6.64	5.54	5.93	5.72	5.63								
NTZRP3003642	10.88	8.03	6.37	13.82	13.96	17.20	12.37	12.40	16.41	*	+	*	+				
NTZRP3003645	4.17	3.53	5.50	3.78	5.31	6.65	5.06	5.99	4.7	**	+						
NTZRP3003648	3.24	3.31	3.16	4.15	4.43	3.91	5.07	3.21	3.18	**	+						
NTZRP3003649	1.14	1.88	2.86	2.19	4.90	3.66	0.71	3.92	1.08								
NTZRP3003659	8.11	4.45	2.20	4.63	4.76	4.42	3.39	3.56	3.36								
NTZRP3003656	5.22	3.74	1.88	3.30	4.87	4.62	3.45	3.40	2.71								
NTZRP3003657	7.45	4.72	4.52	4.36	6.73	4.25	3.17	3.21	3.03								
NTZRP3003662	9.17	7.44	5.08	10.50	15.08	10.64	9.44	8.85	8.35								
NTZRP3003664	8.73	4.21	6.55	11.31	14.75	9.76	9.24	8.50	9.85								
NTZRP3003665	1.46	2.31	3.07	2.00	3.32	2.55	1.63	2.93	1.01								
NTZRP3003671	3.15	3.24	2.25	2.59	7.96	5.47	2.17	4.14	1.93								
NTZRP3003672	4.15	3.09	2.96	4.72	3.77	5.47	2.79	4.69	3.17	*	+						
NTZRP3003673	4.51	3.32	1.35	5.41	6.14	2.58	4.36	4.67	3.13								
NTZRP3003679	24.38	42.38	35.15	32.46	39.83	37.84	41.64	35.07	42.5								
NTZRP3003680	6.95	3.40	1.56	4.84	3.86	4.38	2.61	3.70	3.96								
NTZRP3003686	5.14	3.55	2.82	3.79	4.38	5.04	4.26	3.62	2.84								
NTZRP3003689	3.80	2.46	2.57	6.17	7.73	5.84	3.57	4.94	3.47	**	+						
NTZRP3003697	1.80	2.24	1.34	1.76	2.19	2.72	2.08	3.11	1.51								
NTZRP3003701	1.92	1.12	1.36	1.56	1.36	1.59	2.02	2.99	1.34								
NTZRP3003704	5.17	3.39	3.77	6.61	6.98	7.53	4.92	5.10	3.69	**	+						
NTZRP3003714	3.30	1.91	1.74	4.60	3.93	3.09	3.44	3.54	1.64								
NTZRP3003716	2.44	2.40	1.34	4.13	2.42	3.98	2.31	2.88	2.92								
NTZRP3003721	4.90	3.12	2.28	4.84	6.16	4.98	4.29	3.50	4.65								
NTZRP3003722	8.02	5.81	5.39	6.08	4.24	4.20	2.09	3.67	2.7	*	-						
NTZRP3003726	6.59	6.25	3.44	4.38	3.30	5.00	5.53	4.21	4.73								

1697

1698

NT2RF3003729	3.69	2.88	2.55	4.06	4.92	3.98	2.8	3.60	3.35*	+		
NT2RF3003731	6.61	4.33	5.75	7.10	14.90	8.06	5.99	7.15	5.75			
NT2RF3003740	4.78	3.50	4.29	5.32	3.89	4.79	4.16	4.89	3.61			
NT2RF3003746	5.36	3.49	2.71	5.20	7.52	3.17	3.94	3.31	4.02			
NT2RF3003749	0.76	0.62	0.17	0.29	1.19	1.12	0.64	1.30	0.75			
NT2RF3003754	5.00	3.26	5.25	7.46	7.69	6.19	5.66	4.91	4.55*	+		
NT2RF3003759	1.70	0.69	0.73	1.39	1.06	0.48	0.73	2.09	2.41			
NT2RF3003764	7.97	5.68	5.63	6.40	8.69	7.67	5.36	5.99	4.9			
NT2RF3003766	4.56	2.73	2.99	3.97	4.19	3.87	3.96	3.75	3.32			
NT2RF3003767	6.96	5.70	6.63	13.57	9.41	11.81	7.79	9.76	8.37*	+	*	+
NT2RF3003778	5.19	3.99	4.33	9.90	11.58	8.75	5.62	5.86	5.15**	+		
NT2RF3003779	13.01	5.97	4.99	6.05	7.93	6.85	7.17	5.72	8.58			
NT2RF3003783	19.26	10.08	8.20	11.73	11.20	13.62	12.33	9.52	7.82			
NT2RF3003787	4.90	2.40	2.22	2.44	3.52	4.85	2.78	3.53	7.22			
NT2RF3003789	5.36	4.73	2.56	3.44	7.01	5.23	5.4	5.53	4.62			
NT2RF3003795	2.17	1.85	1.40	3.14	2.08	3.57	2.46	3.18	2.41			
NT2RF3003799	2.89	2.29	1.32	1.87	1.75	2.53	1.45	2.24	2.66			
NT2RF3003800	3.51	2.88	4.22	3.79	5.81	4.55	3.66	3.45	2.49			
NT2RF3003805	6.47	3.37	3.41	4.89	4.12	5.73	3.59	4.60	4.09			
NT2RF3003809	5.03	1.78	2.92	4.79	3.39	3.28	1.85	3.89	3.58			
NT2RF3003819	20.93	12.43	10.20	22.69	23.35	18.68	16.05	13.33	11.82			
NT2RF3003824	12.10	8.20	9.56	14.53	12.56	14.16	10.06	10.73	7.38*	+		
NT2RF3003825	22.51	14.11	14.65	13.44	18.74	15.00	10.89	9.86	10.89			
NT2RF3003828	3.66	3.06	2.75	5.51	4.72	4.12	2.65	4.12	4.14*	+		
NT2RF3003831	2.13	2.74	2.94	4.32	4.71	5.94	3.1	4.50	4.33*	+		
NT2RF3003833	5.17	2.54	2.51	3.72	3.00	5.07	4.52	4.42	4			
NT2RF3003836	7.43	5.49	5.12	9.64	6.79	8.16	7.54	6.97	9.43			
NT2RF3003842	17.19	8.40	7.68	16.76	16.34	13.12	12.09	8.43	8.61			
NT2RF3003843	11.40	7.90	6.65	20.59	22.28	19.09	11.26	10.84	11.37**	+		
NT2RF3003844	12.70	8.55	6.42	7.70	6.74	8.49	13.96	12.46	12.2			
NT2RF3003846	3.76	1.97	2.48	4.49	3.48	4.92	2.73	3.31	3.38			
NT2RF3003849	4.75	3.02	2.95	4.08	4.65	4.41	2.89	4.41	5.12			
NT2RF3003863	8.19	5.27	4.97	5.73	7.14	6.59	9.21	6.75	9.43			
NT2RF3003879	8.87	6.42	4.81	9.09	8.35	8.66	8.21	7.03	8.25			
NT2RF3003874	4.83	4.91	4.32	6.66	5.96	5.92	4.88	5.78	3.78**	+		
NT2RF3003876	8.40	4.71	3.53	8.21	6.66	5.04	3.88	4.35	5.13			
NT2RF3003889	3.42	3.11	2.28	6.01	6.99	4.51	4.71	5.26	4.07*	+	*	+
NT2RF3003889	1.46	1.88	0.92	1.03	3.20	2.06	0.85	2.31	2.72			
NT2RF3003891	1.54	2.90	0.87	1.75	2.99	2.00	1.08	2.80	2.25			
NT2RF3003914	7.95	4.51	4.21	5.57	7.65	7.02	5.69	6.39	7.2			
NT2RF3003915	1.86	2.20	1.19	1.63	2.60	2.36	2.19	3.03	2.1			
NT2RF3003918	5.05	3.66	2.14	2.83	4.62	2.98	3.63	5.42	5.25			
NT2RF3003928	4.98	4.36	2.71	6.50	6.25	5.72	5.51	6.91	3.85*	+		
NT2RF3003924	6.49	3.55	2.01	7.69	8.02	5.14	4.31	3.95	7.6			
NT2RF3003932	3.65	2.42	1.71	4.82	7.98	3.41	2.85	3.76	4.41			
NT2RF3003939	2.69	1.67	1.95	3.86	3.92	3.18	2.41	3.31	2.98*	+		
NT2RF3003940	15.51	8.52	7.81	11.47	11.25	8.35	8.68	9.97	7.23			
NT2RF3003943	3.63	3.38	2.60	2.90	3.77	1.83	2.48	3.48	4.35			
NT2RF3003959	2.34	2.12	1.61	3.04	4.84	3.82	2.42	3.16	4.93*	+		
NT2RF3003963	6.98	5.43	4.54	7.42	7.40	5.93	6.05	7.92	6.84			
NT2RF3003965	44.37	24.77	31.74	35.84	34.50	26.36	12.7	12.05	15.09		*	-
NT2RF3003972	14.33	10.15	6.83	27.44	20.29	23.76	23.62	15.59	17.39*	+		
NT2RF3003973	8.15	5.02	3.70	7.18	5.27	4.94	5.3	5.61	3.97			
NT2RF3003977	11.32	8.28	4.38	9.43	15.88	13.30	10.9	7.26	6.82			
NT2RF3003989	10.84	7.99	7.63	8.16	9.43	9.50	5.75	7.95	4.2			
NT2RF3003982	1.33	3.01	1.15	1.21	2.15	2.04	1.21	3.30	0.58			

111/180

【0752】

【表281】

1699

1700

NT2RF3004009	2.69	2.90	1.66	1.97	4.23	17.93	2.15	5.56	2.09										
NT2RF3004010	4.45	3.19	2.09	6.85	5.45	5.48	2.46	5.01	2.52	*	+								
NT2RF3004011	2.21	2.96	1.05	1.76	3.78	2.06	4.87	2.99	3.16										
NT2RF3004012	10.03	7.36	4.34	11.63	8.96	9.72	6.39	7.58	6.18										
NT2RF3004013	2.84	1.39	1.85	4.23	3.15	3.89	6.12	4.26	2	*	+								
NT2RF3004014	12.35	8.49	6.06	13.19	14.63	10.33	6.81	8.18	5.23										
NT2RF3004015	4.50	2.25	1.85	4.36	3.71	4.81	2.81	2.48	3.43										
NT2RF3004016	4.30	3.53	3.53	4.99	6.65	6.46	4.38	6.03	4.27	*	+								
NT2RF3004017	22.90	14.65	17.74	29.69	32.04	29.24	22.51	18.90	19.9	**	+								
NT2RF3004018	2.52	1.89	2.73	9.78	7.34	7.80	4.71	4.38	4.76	**	+	**	+						
NT2RF3004019	14.33	10.61	5.39	8.88	10.41	10.70	11.54	9.64	11.73										
NT2RF3004020	21.83	11.12	9.61	8.22	9.50	8.35	6.17	5.06	6.39										
NT2RF3004021	10.03	6.48	4.32	11.50	10.92	8.70	7.09	5.39	5.97										
NT2RF3004022	8.89	3.73	4.41	8.80	8.69	8.41	6.86	4.66	5.92										
NT2RF3004023	30.17	20.41	22.51	39.10	49.24	42.11	31.51	31.71	33.47	*	+								
NT2RF3004024	4.37	1.71	1.44	3.41	6.47	4.74	2.67	3.05	2.47										
NT2RF3004025	4.35	3.84	2.26	4.57	5.40	6.36	4.38	3.95	3.58										
NT2RF3004026	3.19	5.38	4.25	5.25	3.73	4.82	2.48	4.55	2.33										
NT2RF3004027	20.57	6.61	6.47	9.24	9.55	7.82	8.89	7.62	7.01										
NT2RF3004028	5.14	4.09	2.46	6.73	5.56	5.86	3.96	3.22	4.36										
NT2RF3004029	4.89	3.98	3.09	4.61	4.46	5.82	3.77	3.33	3.83										
NT2RF3004030	6.60	3.72	3.12	5.82	6.46	5.79	5.42	4.95	4.97										
NT2RF3004031	2.32	2.07	2.04	35.55	41.35	31.65	20.9	19.75	24.51	**	+	**	+						
NT2RF3004032	4.82	3.89	2.80	2.32	2.21	5.07	2.3	4.34	3.24										
NT2RF3004033	6.30	4.80	3.92	7.31	7.31	7.55	5.02	5.55	6.07	*	+								
NT2RF3004034	3.22	2.13	1.57	4.35	5.08	3.83	3.16	6.01	4.35	*	+								
NT2RF3004035	5.89	4.55	3.16	7.72	8.34	6.85	6.58	5.64	6.63	*	+								
NT2RF3004036	14.57	8.24	7.88	13.27	13.82	13.04	10.11	8.74	11.47										
NT2RF3004037	11.19	6.90	6.93	9.17	11.74	10.70	9.42	7.28	9.35										
NT2RF3004038	34.95	22.41	23.25	26.04	28.26	24.02	16.77	18.06	22.74										
NT2RF3004039	6.91	5.16	5.08	8.05	6.96	6.49	5.73	4.85	4.73										
NT2RF3004040	14.03	10.35	8.98	14.12	16.80	14.86	13.91	11.06	10.62										
NT2RF3004041	3.44	1.56	2.05	2.41	2.99	3.58	2.35	2.48	1.77										
NT2RF3004042	3.67	2.75	3.57	6.28	6.18	5.89	7.37	7.97	5.85	**	+	**	+						
NT2RF3004043	8.07	5.45	4.56	6.17	4.28	5.72	6.99	6.13	6.19										
NT2RF3004044	6.56	4.08	2.26	3.88	4.54	4.28	2.91	4.84	3.57										
NT2RF3004045	7.79	6.05	5.54	5.61	5.84	7.93	7.7	7.31	5.13										
NT2RF3004046	3.99	4.60	2.60	5.64	5.29	6.17	3.4	3.66	2.7	*	+								
NT2RF3004047	9.52	6.71	6.33	12.69	13.98	12.98	6.82	6.51	5.79	**	+								
NT2RF3004048	4.17	3.60	3.22	5.35	6.25	6.22	3.75	3.01	3.75	**	+								
NT2RF3004049	2.33	0.68	1.31	1.91	1.20	2.96	1.8	2.34	1.86										
NT2RF3004050	8.37	4.08	5.91	11.26	11.20	6.76	4.54	7.20	6.27										
NT2RF3004051	14.04	5.66	6.06	7.02	12.29	6.24	4.85	4.58	5.6										
NT2RF3004052	11.54	5.42	6.63	7.75	12.77	11.72	5.49	4.47	5.81										
NT2RF3004053	10.44	9.83	8.83	14.00	14.26	11.80	12.41	10.04	10.36	*	+								
NT2RF3004054	2.35	2.27	2.03	3.51	4.57	3.29	3.6	3.97	3.67	*	+	*	+						
NT2RF3004055	10.83	6.54	6.41	8.47	10.58	6.84	7.02	6.54	6.67										
NT2RF3004056	3.85	2.53	2.95	2.95	3.06	3.06	4.12	2.99	2.57										
NT2RF3004057	4.95	2.79	3.03	4.73	4.14	4.86	4.28	4.10	5.09										
NT2RF3004058	4.91	2.40	2.89	6.87	6.50	4.96	4.96	5.23	4.63	*	+								
NT2RF3004059	3.55	2.78	2.14	12.42	8.20	7.94	3.86	4.27	5.18	*	+	*	+						
NT2RF3004060	16.93	6.45	7.83	7.64	9.11	7.10	7.36	6.55	7.25										
NT2RF3004061	5.13	4.26	3.60	4.45	5.10	4.52	4.84	3.47	2.95										
NT2RF3004062	4.82	4.45	3.64	5.22	7.08	6.18	4.56	5.99	5.39	*	+								
NT2RF3004063	1.98	2.17	2.49	2.39	1.93	2.99	2.38	3.72	5.59										
NT2RF3004064	11.77	7.63	9.50	10.32	13.55	13.92	4.51	6.56	5.46			*	-						

112/180

【0753】

【表282】

1702

NTZRP3004263	4.35	2.96	2.85	2.71	3.57	4.45	4.01	4.72	3.41										
NTZRP3004275	3.72	3.04	2.37	3.29	3.02	3.38	3.39	4.75	1.04										
NTZRP3004283	12.87	5.01	5.72	9.16	11.91	6.32	7.38	7.58	6.01										
NTZRP3004289	3.01	2.85	1.46	6.88	5.77	3.72	2.33	3.31	3.68	*									
NTZRP3004294	7.18	34.1	2.73	24.45	29.15	28.18	20.58	15.67	20.34	**									
NTZRP3004299	7.07	5.08	3.77	5.00	5.97	6.16	6.4	6.06	5.61										
NTZRP3004309	10.96	7.28	6.61	7.01	8.68	7.42	5.52	6.85	6.57										
NTZRP3004321	11.18	6.12	7.27	9.56	8.71	10.32	7.19	8.23	10.39										
NTZRP3004322	3.28	2.42	1.89	3.12	2.58	3.70	3.77	3.09	3.39										
NTZRP3004333	6.32	6.72	6.36	11.24	8.54	10.03	4.86	8.82	5.48	*									
NTZRP3004334	4.49	2.34	2.27	5.43	4.10	3.66	2.44	1.92	2.32										
NTZRP3004336	5.86	3.72	2.08	6.83	9.08	6.19	5.13	6.87	5.49										
NTZRP3004338	11.56	5.52	9.71	8.36	5.67	6.93	5.31	4.61	6.32										
NTZRP3004341	2.24	1.74	1.67	2.56	2.48	3.60	1.13	2.33	3.45										
NTZRP3004345	3.27	3.23	2.25	3.71	4.02	3.88	3.2	3.07	3.38										
NTZRP3004348	8.53	5.32	6.83	14.49	13.97	11.82	7.76	7.80	9.23	**									
NTZRP3004349	10.23	7.24	8.20	12.70	11.94	13.01	6.98	7.06	5.47	*									
NTZRP3004355	6.08	5.70	3.65	5.80	6.46	7.00	4.88	5.01	4.97										
NTZRP3004356	13.62	7.29	6.71	12.35	15.04	10.32	9.71	9.44	9.13										
NTZRP3004360	7.52	3.61	3.49	4.81	4.94	4.08	2.07	3.17	4.82										
NTZRP3004361	16.01	7.31	5.66	15.99	14.58	14.13	4.38	5.01	4.13										
NTZRP3004374	7.91	4.13	3.84	7.91	7.91	7.64	5.99	5.39	5.89										
NTZRP3004378	26.21	17.19	14.59	10.81	12.69	11.18	6.13	10.86	9.07										
NTZRP3004399	2.04	2.65	1.39	1.42	2.99	2.67	1.58	2.38	2.75										
NTZRP3004405	3.95	3.77	2.00	4.63	7.05	3.79	3.22	5.96	4.47										
NTZRP3004406	7.20	4.61	5.55	5.61	8.40	5.80	5.82	7.89	6.47										
NTZRP3004411	7.77	3.85	3.09	16.41	12.18	7.61	7.04	7.47	10.13										
NTZRP3004424	4.60	1.42	1.67	3.96	3.79	2.00	1.27	3.09	4.78										
NTZRP3004428	7.15	4.01	3.24	6.42	5.85	3.58	6.97	6.90	7.98										
NTZRP3004432	3.82	2.57	0.97	7.56	9.25	7.81	7.72	10.80	9.98	**									
NTZRP3004434	9.49	5.09	3.75	6.31	8.59	6.98	5.23	4.83	5.64										
NTZRP3004446	6.23	5.35	3.39	6.60	5.96	4.57	2.58	3.47	4.71										
NTZRP3004451	3.49	1.02	1.26	4.53	6.79	4.04	2.13	3.69	4.46										
NTZRP3004454	3.00	1.25	1.36	2.36	2.23	1.93	1.66	2.42	2.5										
NTZRP3004456	16.12	6.82	7.66	12.66	11.01	12.33	11.52	8.75	10.08										
NTZRP3004470	8.70	6.33	3.18	11.68	12.19	10.86	7.44	7.38	5.56	*									
NTZRP3004472	1.89	2.60	1.02	4.08	3.19	3.82	2.45	1.91	1.78	*									
NTZRP3004475	4.99	3.80	4.98	4.54	5.61	3.71	4.53	5.07	4.35										
NTZRP3004480	7.65	5.39	5.29	15.02	14.38	12.51	8.01	7.48	6.29	**									
NTZRP3004481	4.24	6.01	3.44	3.84	4.84	6.10	5.51	4.88	3.41										
NTZRP3004490	1.09	1.00	1.30	1.59	2.17	1.90	1.13	0.94	0.16	*									
NTZRP3004496	11.99	5.64	6.80	14.82	15.35	7.87	12.41	15.48	10.73										
NTZRP3004498	10.57	6.90	5.91	5.39	8.13	7.76	7.22	4.55	5.58										
NTZRP3004500	8.32	3.77	4.24	17.06	17.79	15.82	8.93	7.92	6.72	**									
NTZRP3004504	16.66	9.32	8.13	4.90	5.37	6.99	5.11	6.36	4.24										
NTZRP3004505	8.72	5.28	4.61	4.26	5.67	7.97	8.11	8.94	7.62										
NTZRP3004507	4.86	3.25	3.44	5.31	4.59	4.43	2.27	3.02	3.06										
NTZRP3004519	3.79	1.12	1.28	2.61	2.20	3.15	1.55	1.99	1.88										
NTZRP3004524	1.80	1.60	2.36	2.58	1.67	4.26	2.22	2.30	1.3										
NTZRP3004527	1.16	0.95	0.83	1.29	0.90	1.98	0.25	0.54	0.6										
NTZRP3004534	5.79	3.52	3.93	3.26	4.89	7.19	3.48	2.92	3										
NTZRP3004539	14.05	8.61	6.22	8.74	9.46	10.33	9.99	6.77	9.38										
NTZRP3004541	4.42	3.07	2.91	2.08	4.14	2.58	3.82	4.01	3.83										
NTZRP3004544	2.72	3.68	2.33	4.38	6.86	7.17	4.05	5.81	5.64										
NTZRP3004551	3.07	2.54	2.87	4.53	5.33	3.79	4.49	4.42	3.29	*									
NTZRP3004552	11.09	5.45	4.69	3.94	6.09	8.11	5.03	5.45	2.94										

113/180

1704

N172RP3004557	9.04	5.56	6.56	5.65	4.56	3.38	5.82	5.13	3.59								
N172RP3004561	5.68	3.44	3.35	5.27	5.92	3.88	4.61	5.03	4.06								
N172RP3004566	6.63	6.29	6.33	12.53	11.01	9.77	7.43	8.46	13.57	**	+						
N172RP3004569	6.44	5.29	4.60	10.37	11.99	10.11	4.46	4.55	4.39	**	+						
N172RP3004572	3.83	3.21	2.73	4.62	5.78	5.28	4.28	4.30	2.97	*							
N172RP3004578	3.21	3.44	2.27	5.01	7.11	5.48	3.71	3.96	4.42								
N172RP3004584	3.59	3.64	3.56	3.31	4.74	4.86	3.85	3.43	4.22								
N172RP3004588	3.87	2.70	2.67	8.15	6.21	6.68	4.64	5.48	4.37	**	+		+	+			
N172RP3004594	7.86	6.82	6.37	5.22	4.81	5.30	4.15	4.02	2.13	*	-	*					
N172RP3004603	60.30	35.19	34.71	45.07	50.01	29.71	17.9	21.98	18.08	*	-	*					
N172RP3004612	6.20	3.05	3.45	4.40	4.92	2.76	4.05	3.99	3.11								
N172RP3004617	3.07	2.70	1.70	1.60	2.01	3.22	2.53	2.44	1.96								
N172RP3004618	3.95	2.90	2.07	5.51	5.52	3.64	3.14	3.14	4.18								
N172RP3004625	5.48	4.10	2.95	5.75	7.50	5.56	7.41	6.90	5.44								
N172RP3004635	4.31	4.50	4.46	4.30	6.48	5.74	5.58	3.86	3.99								
N172RP3004640	3.88	3.08	3.28	7.49	7.45	6.73	5.96	5.47	4.27	**	+		+	+			
N172RP3004642	10.28	8.51	8.84	14.09	13.53	15.70	10	10.58	5.55	**	+						
N172RP3004647	7.16	4.79	5.37	9.93	6.54	8.91	7.81	5.99	5.6								
N172RP3004652	9.07	6.60	3.76	13.15	12.30	9.92	7.24	7.33	3.44	*	+						
N172RP3004659	8.16	5.80	4.33	5.00	7.93	7.74	5.7	5.73	5.33								
N172RP3004670	14.41	12.39	9.32	16.29	20.04	15.04	13.36	13.59	15.01								
N172RP3004688	15.39	10.91	11.09	13.50	10.87	9.28	9.4	8.75	8.85								
N172RP30046918	9.99	5.44	8.54	9.01	5.02	7.90	7.84	6.47	7.74								
N172RP30046923	5.20	4.00	3.38	3.86	2.64	2.61	3.51	3.32	2.67								
N172RP30046925	5.36	5.89	4.96	8.91	15.04	11.95	12.96	16.75	13.7	*	+		+	+			
N172RP30046935	8.26	5.47	5.42	13.88	11.54	12.72	5.97	11.43	5.65	**	+						
N172RP30046941	8.69	5.46	1.79	1.69	4.25	2.76	4.28	5.58	4.99								
N172RP3004699	4.05	2.09	2.36	3.68	4.19	3.53	5.9	5.73	3.33								
N172RP30046990	3.62	2.75	1.71	2.29	3.50	3.25	3.01	3.38	3.14								
N172RP30046991	7.84	3.90	4.64	5.71	7.58	5.48	5.27	5.15	5.15								
N172RP30046993	4.66	2.43	2.44	3.26	2.94	4.77	3.68	5.96	2.61								
N172RP30046965	4.21	2.76	2.69	4.09	3.65	3.77	3.32	3.08	2.24								
N172RP30046970	3.16	2.60	2.02	6.63	8.48	9.49	3.2	4.92	3.34	**	+						
N172RP30046974	1.23	0.65	0.45	1.09	0.95	1.43	1.92	3.35	1.24								
N172RP30046978	19.45	8.95	8.65	15.20	11.49	10.74	9.98	6.63	6.98								
N172RP30046980	16.31	10.55	9.31	16.83	24.18	15.57	14.36	10.43	16.69								
N172RP30046989	48.25	34.98	34.96	222.14	203.11	165.35	108.22	86.72	64.03	**	+		+	+			
N172RP300469102	1.59	3.03	0.75	2.02	3.06	3.50	2.33	2.26	2.57								
N172RP300469103	2.96	1.87	1.69	2.51	4.74	2.46	2.75	4.73	2.41								
N172RP300469108	7.32	4.36	4.82	47.03	44.25	37.96	49.26	38.51	49.37	**	+		**	+			
N172RP300469109	12.97	8.34	8.98	9.50	12.20	12.85	13.79	10.89	9.27								
N172RP300469111	1.66	4.14	1.76	3.30	2.22	1.71	2.22	1.42	3.11								
N172RP300469112	12.62	5.96	5.20	13.14	12.78	6.27	9.14	9.28	9.92								
N172RP300469115	6.69	4.45	3.10	4.28	5.71	3.35	6.12	5.23	4.95								
N172RP300469129	5.85	2.83	2.20	2.80	3.92	3.49	3.8	3.85	2.88								
N172RP300469137	6.85	6.38	5.53	4.82	7.68	8.16	4.3	6.03	5.81								
N172RP300469138	31.16	22.51	24.42	13.11	12.17	10.03	14.81	14.41	15.27	**	-	*					
N172RP300469141	4.89	2.65	2.93	4.06	3.52	4.29	2.76	4.18	2.03								
N172RP300469147	2.17	1.29	1.74	2.55	2.46	3.03	2.68	3.29	2.54	*	+		+	+			
N172RP300469150	7.08	4.20	5.06	8.60	7.56	6.25	7.64	8.70	6.48								
N172RP300469151	7.65	4.77	3.15	5.40	5.42	4.70	5.71	4.77	7.73								
N172RP300469157	47.42	28.18	24.63	140.24	151.70	90.24	64.55	61.24	48.04	**	+		+	+			
N172RP300469159	2.50	1.76	1.15	1.15	1.62	2.34	1.61	2.61	1.83								
N172RP300469163	26.39	20.86	16.59	7.91	9.36	8.09	5.61	5.24	4.41	*	-	*		**			
N172RP300469167	3.26	3.04	2.67	3.80	3.99	4.24	2.64	3.85	3.17	**	+						
N172RP300469171	7.53	5.74	5.41	5.89	7.46	4.62	5.54	5.19	6.82								

1705

1706

NTZRP4000178	26.66	17.23	19.20	12.23	15.62	11.17	16.22	18.62	19.97				
NTZRP4000180	17.71	15.54	16.60	7.75	7.76	10.71	9.21	10.11	9.68	**	-	++	-
NTZRP4000183	14.57	9.35	5.99	12.31	15.65	9.34	8.25	9.02	7.47				
NTZRP4000192	9.26	5.09	4.80	6.32	4.48	3.65	4.83	4.74	4.23				
NTZRP4000194	3.63	2.75	1.83	3.79	5.80	2.67	3.51	4.32	4.95				
NTZRP4000196	8.18	4.81	3.10	7.96	7.13	5.03	5.27	5.97	5.49				
NTZRP4000210	28.53	18.46	17.26	28.09	37.05	27.38	24.22	22.19	25.95				
NTZRP4000212	12.06	7.92	6.39	16.76	20.50	16.60	12.99	12.83	12.92	*	+		
NTZRP4000214	10.71	7.74	6.94	13.03	16.29	15.56	10.1	11.28	8.66	*	+		
NTZRP4000216	5.44	4.53	4.98	6.46	9.49	6.90	5.75	6.76	4.95				
NTZRP4000218	7.33	2.22	2.67	4.98	4.33	3.81	3.81	4.58	6.27				
NTZRP4000233	19.92	13.17	10.28	22.13	21.62	13.05	22.62	26.76	25.86			*	+
NTZRP4000243	13.18	9.89	7.93	15.15	23.34	10.85	12.84	16.56	15.03				
NTZRP4000246	33.96	22.95	19.51	28.17	27.99	24.14	21.88	39.67	28.61				
NTZRP4000258	7.99	6.43	5.04	12.08	14.24	11.05	12.85	25.59	17.99	**	+	*	+
NTZRP4000256	2.39	2.62	1.51	3.73	3.59	2.62	3.4	5.63	3.02				
NTZRP4000257	47.78	28.06	32.52	17.19	17.58	12.15	20.3	21.14	18.74	*	-		+
NTZRP4000259	4.57	3.53	4.63	12.50	13.85	8.56	9.95	10.96	10.32	*	+	**	+
NTZRP4000261	4.69	3.90	2.69	4.69	4.12	2.59	6.07	3.27	3.23				
NTZRP4000262	8.40	4.25	5.05	10.81	7.69	5.18	7.05	4.76	3.22				
NTZRP4000263	2.39	2.26	1.46	3.24	1.78	2.52	2.31	2.43	1.67				
NTZRP4000280	19.84	10.94	16.02	14.51	20.53	17.86	16.38	15.79	14.33				
NTZRP4000286	14.05	12.14	5.20	8.66	7.23	8.18	6.73	10.62	6.93				
NTZRP4000290	4.20	3.07	2.79	5.43	3.58	4.59	3.38	3.10	2.4				
NTZRP4000291	18.51	15.32	18.47	45.30	38.54	34.77	17.5	19.25	13.11	**	+		
NTZRP4000301	2.59	1.81	1.04	2.23	2.98	3.54	2.54	3.49	1.63				
NTZRP4000312	4.56	1.79	4.33	4.54	4.75	3.56	5.14	2.41	5.06				
NTZRP4000331	13.60	6.74	4.54	13.92	11.99	10.85	8.51	8.80	9.62				
NTZRP4000323	3.58	2.53	1.59	2.86	3.50	3.23	2.71	3.60	1.23				
NTZRP4000334	7.25	5.08	2.70	5.19	6.35	3.74	5.48	4.98	4				
NTZRP4000334	13.97	11.43	12.75	30.03	27.15	21.64	10.28	10.30	9.71	**	+	*	-
NTZRP4000343	4.98	3.25	2.65	4.86	5.56	3.68	3.76	4.39	3.15				
NTZRP4000348	3.02	1.79	1.77	4.45	3.35	4.09	4.17	3.46	2.74	*	+		
NTZRP4000349	2.02	3.31	1.01	2.05	0.64	3.58	0.41	1.43	0.27				
NTZRP4000355	10.07	4.28	4.14	7.89	8.66	7.17	5.76	4.78	6.28				
NTZRP4000356	10.81	5.71	5.12	9.75	8.69	6.70	12.73	12.78	15.8			*	+
NTZRP4000368	5.76	3.41	2.25	11.67	15.48	9.10	8.87	7.21	7.44	*	+	*	+
NTZRP4000367	2.23	2.01	1.13	1.88	2.90	1.83	2.17	1.67	2.44				
NTZRP4000370	4.54	3.75	1.61	3.50	4.39	3.20	3.15	3.31	3.03				
NTZRP4000373	4.40	4.53	4.20	4.85	4.38	4.02	3.74	3.46	2.82			*	
NTZRP4000376	3.46	3.35	3.32	5.35	3.36	3.31	2.76	4.60	2.39				
NTZRP4000381	3.20	2.91	2.81	7.76	5.97	5.48	3.69	3.62	2.58	**	+		
NTZRP4000388	507.68	363.39	334.24	288.84	217.90	196.35	431.3	437.24	362.7				
NTZRP4000390	19.01	14.68	11.68	24.99	29.51	23.19	15.68	13.59	14.64	*	+		
NTZRP4000393	3.40	2.87	1.85	2.59	3.15	3.33	5.06	3.98	3.29				
NTZRP4000398	5.34	4.25	2.50	10.36	14.48	10.01	6.8	5.94	5.69	*	+		
NTZRP4000406	9.30	5.25	6.26	5.59	5.04	6.35	7.54	6.32	4.52				
NTZRP4000407	5.98	4.41	3.78	8.29	7.16	4.70	4.32	5.68	5.13				
NTZRP4000413	1.40	1.18	0.62	0.72	1.57	3.58	1.37	2.49	1.36				
NTZRP4000415	10.74	4.75	5.55	8.27	6.74	8.60	4.84	5.48	2.05				
NTZRP4000417	7.49	5.67	3.62	5.24	6.05	4.58	5.78	5.18	6.52				
NTZRP4000423	10.91	8.43	6.08	17.00	12.75	12.74	5.48	6.12	5.86	*	+		
NTZRP4000424	4.48	2.86	1.81	7.46	7.77	6.37	5.69	7.35	4.76	**	+		
NTZRP4000447	13.10	8.03	11.15	9.03	13.44	9.03	6.38	5.33	5.62			*	-
NTZRP4000448	2.34	1.79	0.84	4.19	6.84	6.98	5.24	4.20	3.76	*	+	*	+
NTZRP4000449	2.70	2.01	2.13	2.07	1.89	2.22	2.41	2.65	1.44				

115/180

【0756】

【表285】

1707

1708

NT2RP4000443	7.28	6.16	3.48	2.33	2.43	4.15	1.8	4.72	0.91				
NT2RP4000444	1.01	1.01	1.48	2.29	2.70	1.92	2.22	2.27	0.83	*	+		
NT2RP4000445	13.97	7.10	6.36	13.16	13.46	10.68	8.85	8.11	5.28				
NT2RP4000447	6.68	4.82	2.84	3.69	4.73	3.69	4.6	3.98	5.62				
NT2RP4000461	5.28	3.96	3.32	7.87	8.68	6.42	5.85	6.52	5.36	*	+		
NT2RP4000462	8.07	4.05	4.23	7.49	8.39	11.75	6.93	5.29	4.06				
NT2RP4000463	9.18	6.18	6.85	10.59	9.85	9.05	5.78	4.84	4.27				
NT2RP4000471	3.55	1.94	1.96	3.21	3.41	4.25	4.22	4.59	2.95				
NT2RP4000472	3.05	2.42	1.96	12.20	8.76	6.84	4.36	5.24	4.11	*	+	*	+
NT2RP4000476	1.50	1.02	0.85	12.49	11.85	10.88	21.84	18.65	17.71	**	+	**	+
NT2RP4000480	15.36	6.51	5.30	5.47	9.87	5.81	7.44	7.54	5.87				
NT2RP4000481	3.47	2.35	0.78	2.35	2.92	2.36	3.06	3.89	4.07				
NT2RP4000483	2.86	2.52	1.45	2.10	2.49	1.39	3.11	4.18	2.64				
NT2RP4000487	3.11	1.79	1.56	6.59	4.70	2.73	3.7	3.87	2.46				
NT2RP4000496	0.65	2.01	0.43	0.74	1.20	0.89	1.64	1.30	1.26				
NT2RP4000497	6.68	4.62	5.43	14.85	10.68	12.20	7.76	11.46	5.67	**	+		
NT2RP4000498	4.09	1.89	2.15	3.59	3.39	3.97	3.69	5.45	2.91				
NT2RP4000500	3.65	2.95	1.78	3.44	3.70	2.25	3.4	3.63	2.11				
NT2RP4000507	15.14	8.22	5.69	11.50	10.49	7.06	7.7	7.22	9.04				
NT2RP4000515	15.49	10.59	8.57	12.80	13.50	16.10	12.82	10.19	8.69				
NT2RP4000516	7.24	4.39	3.65	20.66	19.29	17.91	10.11	9.21	8.83	**	+	*	+
NT2RP4000517	3.07	2.43	1.84	4.04	5.74	5.81	3.42	4.89	3.38	*	+		
NT2RP4000518	4.18	1.91	2.39	4.28	2.50	2.78	3.19	3.42	2.91				
NT2RP4000519	1.25	1.47	1.18	2.14	1.80	1.86	1.53	2.34	1.09	**	+		
NT2RP4000524	0.66	1.08	0.33	1.66	1.94	1.79	1.87	1.81	1.62	**	+	**	+
NT2RP4000528	1.96	2.16	0.43	1.52	2.71	2.98	1.9	3.84	1.18				
NT2RP4000537	40.32	18.87	17.18	18.72	15.16	10.99	14.21	10.18	11.8				
NT2RP4000541	6.42	4.52	3.64	6.16	5.27	3.57	5.96	5.32	5.79				
NT2RP4000540	7.15	4.38	3.94	5.71	5.28	6.49	7.13	6.85	7.19				
NT2RP4000545	22.00	12.60	11.90	35.02	30.28	28.43	15.85	15.53	13.71	*	+		
NT2RP4000546	3.49	2.74	2.72	5.16	6.84	5.20	2.65	5.26	4.13	*	+		
NT2RP4000549	10.31	6.26	6.97	10.02	6.99	7.06	17.04	10.70	13.71				
NT2RP4000556	4.79	2.38	2.09	2.96	4.25	3.16	3.01	3.93	2.39				
NT2RP4000557	2.43	1.89	1.59	3.06	2.06	2.13	1.6	1.76	2.34				
NT2RP4000558	7.85	4.61	3.47	5.80	4.60	4.48	8.11	4.97	5.07				
NT2RP4000560	11.62	8.43	5.62	16.38	11.32	8.62	10.3	8.86	6.76				
NT2RP4000568	0.86	1.06	0.72	1.99	2.89	2.56	1.2	1.79	1.98	**	+	*	+
NT2RP4000583	9.91	5.21	4.91	9.30	13.09	14.53	6.79	5.52	7.23				
NT2RP4000585	3.74	2.64	3.88	4.44	2.94	3.43	2.78	2.68	3.99				
NT2RP4000588	1.78	1.61	0.91	2.23	3.68	2.01	2.78	3.01	2.89		**	+	
NT2RP4000590	7.09	4.23	3.81	4.80	5.51	5.49	5.51	5.97	3.62				
NT2RP4000599	1.53	1.26	0.87	1.24	1.41	1.06	0.44	2.70	0.51				
NT2RP4000603	11.90	6.03	3.83	6.61	6.16	3.84	4.98	5.30	6.79				
NT2RP4000607	9.25	5.44	5.52	6.95	7.07	10.29	4.24	5.47	7.66				
NT2RP4000614	18.95	12.78	10.17	25.67	26.47	23.13	9.33	11.19	9.77	*	+		
NT2RP4000634	4.83	2.61	1.81	7.54	6.71	5.97	5.4	7.61	4.39	*	+		
NT2RP4000638	3.55	2.37	1.27	3.88	3.82	3.28	2.34	4.08	2.48				
NT2RP4000648	3.49	3.15	1.64	4.18	4.00	1.87	2.79	3.50	2.8				
NT2RP4000657	7.42	4.66	4.89	3.76	5.89	4.90	4.79	4.73	4.39				
NT2RP4000691	3.57	4.48	4.25	6.09	7.82	5.58	5.65	7.17	5.49	*	+	*	+
NT2RP4000697	11.06	7.17	4.24	7.59	7.47	5.97	4.38	5.33	7.55				
NT2RP4000704	9.94	4.45	4.08	7.72	7.80	6.93	8.9	11.64	11.09				
NT2RP4000710	39.78	22.43	20.25	37.57	42.17	34.47	22.39	29.16	28.71				
NT2RP4000713	3.09	1.40	0.88	3.21	4.08	3.16	3.3	5.18	2.97				
NT2RP4000724	3.53	1.86	1.77	4.48	4.24	3.42	3.25	6.43	3.91				
NT2RP4000725	4.59	2.50	2.14	3.16	3.33	2.21	3.39	4.06	2.51				

116/180

【0757】

【表286】

1709

1710

NT2RP4000728	21.11	12.54	13.41	26.39	33.93	29.91	18.2	20.00	17.52	*	+	
NT2RP4000737	2.29	1.59	0.36	2.95	3.74	3.56	1.99	4.28	1.59	*	+	
NT2RP4000739	3.68	1.68	1.40	3.64	3.60	3.19	3.01	1.32	2.06			
NT2RP4000749	4.61	2.23	2.17	5.43	5.08	3.32	3.77	2.84	2.99			
NT2RP4000769	4.46	2.77	1.61	5.35	5.75	3.05	3.69	3.92	2.49			
NT2RP4000774	7.04	3.62	4.69	6.48	7.03	5.14	4.99	3.77	3.67			
NT2RP4000781	1.78	1.82	2.45	2.48	1.82	2.08	1.95	1.67	1.08			
NT2RP4000783	5.52	3.48	3.60	5.32	4.17	5.29	1.54	2.21	1.91	*	-	
NT2RP4000787	(0.08)	0.27	0.06	0.45	0.09	1.07	0.1	0.13	-0.1			
NT2RP4000788	7.00	4.42	3.89	7.56	7.52	5.50	5.26	4.25	3.66			
NT2RP4000792	9.90	5.45	5.18	4.82	3.85	3.35	2.89	1.10	1.13			*
NT2RP4000809	138.97	85.82	100.50	13.12	12.28	11.89	8.69	10.55	11.51	**	-	**
NT2RP4000817	6.53	3.13	3.81	7.81	8.21	7.10	5.75	6.24	6.14	*	+	*
NT2RP4000821	10.40	5.88	5.97	8.60	9.00	10.24	19.32	14.83	13.61	*	+	*
NT2RP4000822	7.54	4.48	4.61	11.43	10.03	11.32	7.11	5.54	4.78	**	+	*
NT2RP4000823	6.10	4.87	4.52	6.50	4.58	4.69	17.58	17.55	14.17			**
NT2RP4000831	4.53	2.70	1.65	4.00	4.27	4.75	3.68	4.83	3.77			*
NT2RP4000833	9.98	4.61	3.88	12.93	9.95	9.75	7.85	6.14	9.61			*
NT2RP4000837	16.84	7.67	8.19	4.27	7.04	6.55	7.9	6.72	7.63			*
NT2RP4000839	8.09	4.28	3.15	6.64	6.35	8.56	6.01	3.49	4.81			*
NT2RP4000846	7.97	4.70	3.74	7.70	5.83	5.14	6.12	4.09	4.53			*
NT2RP4000848	5.78	2.64	3.11	8.90	6.26	8.65	7.07	7.56	8.46	*	+	*
NT2RP4000855	3.22	3.08	1.54	2.41	2.92	2.82	2.82	2.57	2			*
NT2RP4000863	3.79	2.50	2.36	1.24	1.67	1.78	2	2.70	1.71			*
NT2RP4000865	9.55	7.40	5.94	26.23	26.54	18.52	8.98	8.90	8.56	**	+	*
NT2RP4000873	8.88	4.73	4.97	9.82	9.15	8.69	10.43	4.81	6.51			*
NT2RP4000874	5.60	3.25	3.18	4.02	6.09	6.60	5.15	3.17	5.54			*
NT2RP4000875	10.06	7.69	6.92	10.24	9.60	8.28	5.61	5.34	4.98			*
NT2RP4000878	15.02	8.48	6.31	16.61	14.17	15.37	18.42	13.92	17			*
NT2RP4000879	1.68	0.79	0.77	1.38	2.21	2.54	2.35	2.03	1.86			*
NT2RP4000880	5.88	4.11	3.04	9.39	7.05	7.35	6.97	5.69	5.31	*	+	*
NT2RP4000891	102.85	62.84	77.22	114.50	151.60	104.23	43.98	42.97	34.75			*
NT2RP4000894	8.78	5.12	4.69	6.91	6.62	9.49	7.97	4.83	7.88			*
NT2RP4000898	0.75	1.21	0.33	0.94	1.28	0.69	1.75	1.00	0.58			*
NT2RP4000899	14.91	8.73	9.27	8.87	7.17	6.06	2.92	6.91	6.96			*
NT2RP4000907	7.23	4.77	4.04	8.01	14.43	8.65	11.43	9.68	10.25			**
NT2RP4000908	3.70	3.82	2.81	5.39	5.05	5.27	4.11	5.22	3.41	**	+	*
NT2RP4000918	11.95	5.36	6.97	10.03	8.98	9.73	9.64	9.49	7.69			*
NT2RP4000918	10.45	8.95	8.11	12.80	9.01	11.75	7.94	8.71	6.88			*
NT2RP4000925	1.77	2.18	1.68	2.08	2.56	3.09	1.91	2.37	0.93			*
NT2RP4000927	2.00	0.98	0.64	1.21	1.11	1.91	1.67	2.03	0.45			*
NT2RP4000928	8.63	5.13	3.60	5.86	6.72	6.51	5.18	4.85	6.78			*
NT2RP4000929	1.61	1.10	1.06	1.59	2.36	1.14	0.96	1.23	1.92			*
NT2RP4000946	3.91	2.24	2.26	7.89	6.10	6.89	5.7	5.35	4.43	**	+	*
NT2RP4000947	1.12	1.54	1.05	1.80	1.82	0.62	1.3	1.55	0.89			*
NT2RP4000949	16.12	8.67	10.24	5.88	3.51	5.79	19.02	19.45	15.95			*
NT2RP4000955	9.21	5.55	4.76	5.43	4.34	5.39	4.04	4.48	4.02			*
NT2RP4000959	16.07	16.16	17.01	17.30	15.74	18.65	13.76	14.61	12.03			*
NT2RP4000962	4.28	2.72	4.02	3.76	4.20	2.99	2.02	3.10	1.89			*
NT2RP4000973	6.76	3.78	2.61	4.40	5.06	4.18	8.32	7.27	7.89			*
NT2RP4000975	4.74	2.41	1.77	5.26	4.90	3.72	2.88	4.71	4.07			*
NT2RP4000979	6.80	3.38	3.74	6.77	5.99	3.62	6.11	4.01	4.79			*
NT2RP4000984	3.24	3.46	2.61	2.85	2.49	5.25	1.35	3.81	1.22			*
NT2RP4000986	3.13	2.19	3.27	2.70	3.05	3.24	3.2	4.03	2.69			*
NT2RP4000988	4.24	3.53	3.97	6.52	7.14	6.40	4.03	5.72	2.89	**	+	*
NT2RP4000989	4.55	3.53	3.49	5.18	3.51	4.95	4.91	5.46	4.69	*	+	*

117/180

[0758]

[表287]

1711

1712

NT2RP4000990	0.91	1.17	0.68	5.32	4.83	4.20	3.51	3.92	3.51	**	+	**+
NT2RP4000994	6.03	3.61	2.39	2.73	3.58	3.95	4.94	3.50	5.8			
NT2RP4000996	6.29	4.22	3.37	8.35	8.21	4.36	4.41	5.02	6.24			
NT2RP4000997	61.78	21.49	33.43	48.43	44.30	38.85	25.67	23.78	20.69			
NT2RP4001001	5.72	4.90	3.47	5.67	6.31	7.83	5.36	5.68	6.44			
NT2RP4001004	2.47	1.20	1.29	1.66	1.42	2.31	0.88	2.30	2.26			
NT2RP4001006	6.01	3.42	6.46	5.11	3.94	7.35	4.19	4.92	4.66			
NT2RP4001009	8.55	4.50	6.33	9.69	4.66	6.57	7.89	8.50	7.3			
NT2RP4001010	2.33	1.99	3.31	3.50	2.89	4.49	3.41	2.18	2.22			
NT2RP4001013	24.76	12.16	10.77	11.37	8.47	9.68	9.97	7.96	8.93			
NT2RP4001029	12.87	4.18	5.93	5.61	5.98	4.93	3.75	3.75	3.77			
NT2RP4001036	12.25	7.10	7.56	11.16	11.59	9.83	8.7	8.94	6.61			
NT2RP4001041	12.91	6.26	9.00	10.06	7.34	6.55	5.46	5.13	5.26			
NT2RP4001042	19.25	12.69	10.60	14.77	15.99	12.64	7.69	8.09	6.86			
NT2RP4001046	7.12	4.49	4.11	7.61	7.51	8.81	6.32	5.77	5.49			
NT2RP4001050	2.62	1.51	1.21	2.43	2.08	3.36	1.88	2.76	1.6			
NT2RP4001051	6.34	2.77	3.34	9.61	5.53	9.29	3.29	7.15	4.7			
NT2RP4001057	8.53	5.25	3.91	5.29	5.25	3.31	3.9	2.95	5.29			
NT2RP4001063	10.42	5.01	5.86	6.23	5.90	5.66	7.16	6.14	5.17			
NT2RP4001064	8.38	3.24	3.12	6.83	5.16	4.28	7.84	6.40	8.7			
NT2RP4001067	3.31	1.58	2.32	2.67	2.92	2.43	3.23	3.73	3.54			
NT2RP4001078	4.41	2.35	1.43	3.09	2.15	3.81	2.07	3.77	2.39			
NT2RP4001079	3.33	2.47	3.40	5.24	5.12	4.58	4.51	5.49	6.01	**	+	+
NT2RP4001080	1.87	1.13	0.72	2.09	1.21	1.64	1.8	2.65	1.93			
NT2RP4001086	6.48	4.50	4.95	6.91	6.12	6.66	5.1	5.61	4.86			
NT2RP4001095	9.39	3.28	2.95	11.12	8.02	6.83	6.4	5.11	6.99			
NT2RP4001098	8.66	3.42	3.13	5.99	6.59	3.50	4.06	3.58	3.83			
NT2RP4001100	15.58	6.86	5.99	15.36	16.25	10.53	11.07	8.66	10.12			
NT2RP4001105	11.53	6.11	5.68	11.42	12.40	12.53	6.82	8.59	7.03			
NT2RP4001110	4.14	2.11	2.03	3.53	3.73	5.22	7.74	9.16	5.7	*	+	
NT2RP4001115	8.23	4.76	5.40	7.44	6.61	6.42	6.49	8.54	8.25			
NT2RP4001117	5.86	2.61	3.66	4.84	5.68	5.67	6.82	7.82	11.35			
NT2RP4001122	4.53	2.89	4.44	5.52	5.25	6.14	3.92	5.33	5.22	*	+	
NT2RP4001123	11.03	6.64	4.19	7.23	8.62	6.22	6.52	4.59	7.16			
NT2RP4001126	12.30	8.16	5.35	14.50	10.35	10.40	6.7	7.95	9.08			
NT2RP4001127	2.67	1.52	0.45	2.09	2.22	1.57	1.96	4.17	3			
NT2RP4001138	3.41	2.11	1.63	1.48	2.64	1.74	2.14	3.24	4.17			
NT2RP4001143	6.89	2.21	3.01	4.13	4.68	5.32	4.17	5.67	5.66			
NT2RP4001148	1.94	1.16	1.16	2.70	2.05	0.60	1.41	3.15	1.62			
NT2RP4001149	4.34	2.11	2.80	3.19	3.00	3.41	3.12	4.58	4.05			
NT2RP4001150	4.09	2.84	2.82	5.63	5.48	6.34	4.62	4.61	4.79	**	+	+
NT2RP4001159	8.72	3.82	5.00	5.57	8.96	6.80	7.8	6.33	6.38			
NT2RP4001162	3.97	2.49	1.88	3.46	2.36	3.14	3.98	2.29	2.75			
NT2RP4001170	9.81	5.75	5.29	2.68	3.96	2.23	2.4	2.44	1.3	*	-	
NT2RP4001174	6.78	5.08	5.60	9.49	9.90	7.92	7.08	5.86	4.66	*	+	
NT2RP4001175	19.07	9.74	10.40	16.34	17.86	15.79	8.78	8.58	11.27			
NT2RP4001176	62.90	39.84	55.63	104.65	115.71	110.77	63.62	58.35	46.85	**	+	
NT2RP4001184	10.39	5.65	5.39	5.95	4.48	5.41	4.76	4.78	4.24			
NT2RP4001198	10.79	4.11	5.82	13.69	9.03	11.21	14.64	14.06	13.84	*	+	
NT2RP4001199	2.92	0.71	0.91	2.99	2.97	1.91	3.68	2.25	2.92			
NT2RP4001206	13.96	4.32	7.41	11.41	10.25	10.46	8.73	9.26	10.42			
NT2RP4001207	3.37	2.92	1.08	2.45	1.58	1.84	2.26	2.66	0.61			
NT2RP4001210	2.36	1.47	2.10	3.13	2.39	1.71	1.5	2.49	2.3			
NT2RP4001213	10.44	5.34	6.49	11.64	9.13	13.58	7.15	5.01	5.42			
NT2RP4001214	0.95	1.06	0.59	2.80	1.54	8.36	1.71	2.54	1.49	*	+	
NT2RP4001219	2.55	2.66	2.86	4.42	15.66	4.45	5.58	7.03	5.57	**	+	+

118/180

【0759】

【表288】

1713

1714

NT2RP4001328	6.93	2.54	3.03	5.28	9.41	5.96	8.24	4.93	8.37										
NT2RP4001335	6.11	4.31	3.21	5.70	5.94	5.25	5.94	4.41	5.1										
NT2RP4001336	4.51	1.77	2.22	4.07	5.11	4.94	4.27	3.05	2.43										
NT2RP4001337	6.40	4.02	2.26	5.05	5.34	3.44	5.95	5.21	4.31										
NT2RP4001340	5.39	3.07	4.18	8.97	9.59	5.62	5.8	6.24	6.64			*							
NT2RP4001341	14.65	12.44	12.58	14.19	12.55	13.99	17.34	12.10	15.2										
NT2RP4001374	4.71	4.57	4.07	7.45	6.65	6.76	5.26	6.13	6.26			**	+	*					
NT2RP4001376	15.31	8.46	8.50	10.61	14.38	10.37	11.44	11.39	8.98										
NT2RP4001383	63.21	34.01	32.33	24.21	25.03	19.31	48.06	42.63	46.56										
NT2RP4001399	15.00	9.02	6.78	6.64	8.24	7.13	7.92	6.14	6.14										
NT2RP4001313	3.06	1.56	1.37	2.51	0.89	2.21	1.62	2.23	2.1										
NT2RP4001315	3.67	2.67	2.40	3.93	5.09	3.45	3.89	3.89	4.16										
NT2RP4001320	9.02	4.65	5.15	9.20	8.51	8.68	15.43	12.65	14.49			**		+					
NT2RP4001325	12.74	11.57	11.78	16.64	15.36	9.87	12.12	10.53	7.42										
NT2RP4001336	6.40	4.16	5.13	5.38	3.83	5.19	4.39	4.05	2.52										
NT2RP4001339	3.62	2.24	4.32	4.37	4.09	4.92	3.51	4.78	3.43										
NT2RP4001343	8.44	4.63	3.67	7.94	6.79	5.81	5.7	6.09	6.51										
NT2RP4001344	5.76	3.40	4.09	5.03	5.50	6.54	6.12	6.22	5.58										
NT2RP4001345	6.21	3.12	2.61	3.29	6.07	5.15	4.25	4.33	4.38										
NT2RP4001351	11.92	6.04	5.53	9.86	6.47	8.71	6.54	7.28	6.61										
NT2RP4001353	1.80	1.08	1.42	2.16	2.00	2.04	2.15	2.48	2.23			*		+	*				
NT2RP4001355	2.54	1.08	2.05	2.40	2.01	1.99	2.51	3.62	2.23										
NT2RP4001367	23.22	13.41	17.84	6.30	4.94	5.47	9.28	11.30	7.57			*		*					
NT2RP4001372	5.35	2.77	2.56	3.34	4.53	3.99	4.57	5.24	5.57										
NT2RP4001373	10.60	5.25	4.77	8.11	9.86	9.53	6.1	5.34	6.98										
NT2RP4001375	5.11	3.33	2.60	2.66	4.56	3.81	2.85	3.42	3.31										
NT2RP4001379	3.86	2.14	2.09	2.83	2.70	4.72	3.26	3.43	2.58										
NT2RP4001381	8.37	5.24	5.75	10.66	11.10	10.55	6.09	7.62	6.54			*		+					
NT2RP4001386	3.36	2.18	2.25	6.41	4.78	6.49	3.68	5.89	3.24			**		+					
NT2RP4001389	10.33	5.90	8.63	13.74	8.10	10.59	13.58	10.92	11.95										
NT2RP4001396	1.51	0.17	0.39	1.10	1.45	1.19	1.43	2.48	0.52										
NT2RP4001407	2.74	1.02	1.62	3.87	3.78	1.98	2.72	2.67	1.52										
NT2RP4001409	7.90	3.42	3.68	8.04	5.25	6.06	3.89	2.35	3.87										
NT2RP4001410	41.71	16.67	20.24	29.88	31.04	31.69	28.88	20.00	22.74										
NT2RP4001414	11.73	6.50	5.48	10.69	11.38	10.17	10.68	8.69	10.89										
NT2RP4001424	3.25	2.51	1.43	4.18	3.70	4.01	2.5	5.15	3.66			*		+					
NT2RP4001433	10.93	1.50	1.13	15.16	15.56	3.13	10.41	4.52	7										
NT2RP4001438	8.06	6.23	6.43	14.12	10.57	11.39	6.77	9.65	7.69			*		+					
NT2RP4001442	5.25	2.76	3.72	6.62	2.55	2.88	2.74	3.33	2.46										
NT2RP4001447	1.94	1.07	2.00	4.12	2.36	3.98	1.68	3.22	0.71			*		+					
NT2RP4001446	13.13	5.79	4.82	7.69	5.30	6.70	2.91	4.53	3.9										
NT2RP4001467	4.50	1.22	1.33	0.82	1.55	1.40	3.66	4.13	3.7										
NT2RP4001472	4.77	3.08	3.33	7.29	7.84	10.23	7.79	8.21	9.21			*		+	**				
NT2RP4001474	2.86	1.72	1.90	2.18	3.93	2.05	1.94	3.80	3.06										
NT2RP4001480	2.29	1.49	1.84	3.04	2.50	2.14	2.24	3.68	2.54										
NT2RP4001488	5.16	2.65	2.75	5.33	5.10	5.16	4.15	4.07	6.19										
NT2RP4001492	5.93	3.30	2.87	5.58	3.40	4.66	3.78	4.60	5.29										
NT2RP4001498	2.17	1.63	1.33	2.59	1.07	2.19	2.61	1.92	1.74										
NT2RP4001503	36.00	12.08	15.43	15.15	11.96	14.06	11.15	10.06	10.33										
NT2RP4001503	12.74	6.75	6.97	11.88	9.69	8.87	5.71	4.79	6.02										
NT2RP4001507	5.29	3.53	4.09	6.91	8.58	5.74	3.85	4.47	6.06			*		+					
NT2RP4001510	9.01	6.05	7.69	15.28	11.96	14.90	7.45	6.33	7.03			**		+					
NT2RP4001516	6.51	3.15	3.51	3.57	3.42	3.81	3.63	4.76	4.46										
NT2RP4001528	26.12	11.82	16.11	17.96	13.99	17.71	16.8	15.23	12.31										
NT2RP4001523	3.37	1.82	2.58	4.23	3.77	4.26	2.29	4.88	4.21			*		+					
NT2RP4001524	11.16	7.76	6.79	8.80	7.75	9.91	6.38	9.28	5.14										

119/180

【0760】

【表289】

1715

1716

NT2RP4001529	9.24	4.27	3.42	3.66	4.21	3.95	6.65	3.78	5.28										
NT2RP4001531	7.58	4.22	3.87	4.40	6.79	5.07	4.85	4.25	5.33										
NT2RP4001546	27.96	14.34	13.14	33.50	26.35	22.36	39.72	37.62	29.88										
NT2RP4001547	5.16	3.87	3.59	6.27	5.81	5.41	6.77	5.69	7.74	*									
NT2RP4001551	4.66	2.25	2.91	1.72	2.50	2.23	1.06	2.31	2.02										
NT2RP4001555	2.63	1.70	1.48	1.84	1.34	1.78	3.29	2.29	1.99										
NT2RP4001567	4.17	2.21	3.48	5.17	4.12	2.97	3.53	3.55	4.6										
NT2RP4001568	24.66	11.55	19.71	26.48	16.71	27.97	21.61	20.91	21.83										
NT2RP4001569	13.23	7.51	6.17	8.88	7.94	7.65	6.86	6.56	7.44										
NT2RP4001571	3.88	2.14	1.80	4.74	3.69	4.71	3.97	5.20	7.86										
NT2RP4001574	8.95	4.84	4.26	8.19	9.78	5.65	6.26	6.22	8.16										
NT2RP4001575	8.04	4.77	3.76	6.08	7.50	5.82	4.63	5.56	5.85										
NT2RP4001578	11.18	4.73	6.33	7.50	4.87	4.81	7.41	8.00	7.35										
NT2RP4001592	9.35	5.87	4.90	5.95	6.70	4.56	3.37	8.97	5.41										
NT2RP4001593	6.28	4.83	5.72	9.71	12.44	12.90	7.66	7.56	6.44	**									
NT2RP4001605	4.40	2.61	3.07	7.26	7.76	5.64	5.16	7.35	8.18	*									
NT2RP4001606	13.15	5.10	4.06	9.17	7.65	6.75	3.7	4.31	6.28										
NT2RP4001607	3.47	1.57	1.29	3.76	4.78	2.65	1.67	3.06	4.34										
NT2RP4001610	4.08	2.08	1.47	3.77	3.73	2.68	2.34	4.35	2.92										
NT2RP4001614	2.75	1.07	1.10	2.96	1.97	1.29	2.18	3.56	3.15										
NT2RP4001623	3.08	1.60	1.52	2.58	2.94	2.80	1.24	3.23	2.34										
NT2RP4001626	19.42	15.83	18.19	15.38	17.59	13.04	1.75	4.18	2.95	**									
NT2RP4001634	4.38	2.77	2.43	4.92	4.36	4.52	1.82	3.51	2.53										
NT2RP4001638	2.68	1.70	0.84	1.98	2.75	2.80	1.64	3.48	1.26										
NT2RP4001644	3.61	2.50	2.30	4.35	3.54	2.45	4.35	2.84	4.05										
NT2RP4001646	20.39	11.21	10.21	30.98	19.98	25.17	21.75	14.88	9.56										
NT2RP4001656	6.55	3.72	4.64	5.20	5.23	4.49	4.29	3.23	2.79										
NT2RP4001666	5.11	3.28	3.35	4.54	4.56	3.95	3.53	3.52	3.51										
NT2RP4001670	7.31	3.77	5.28	4.59	6.96	4.67	4.23	4.15	4.55										
NT2RP4001677	16.68	12.12	14.19	29.06	40.57	32.81	33.86	36.13	36.39	**									
NT2RP4001679	11.61	4.52	5.94	19.33	14.25	14.99	8.64	9.90	7.91	*									
NT2RP4001685	20.41	7.98	11.64	19.72	19.63	15.23	7.89	9.75	7.32										
NT2RP4001696	6.64	4.27	3.64	4.33	3.58	5.85	4.75	3.99	3.79										
NT2RP4001699	1.63	1.58	0.71	2.91	1.63	2.15	3.74	2.30	2.42										
NT2RP4001717	3.33	4.49	3.61	5.92	6.26	5.39	5.73	6.49	5.79										
NT2RP4001719	3.81	3.40	2.34	4.26	2.94	3.04	4.14	3.43	2.54										
NT2RP4001725	4.09	3.08	1.88	3.37	4.40	3.86	2.62	4.74	3.15										
NT2RP4001726	4.90	3.18	3.91	4.82	4.39	4.14	4.14	5.24	5.01										
NT2RP4001730	0.78	0.69	0.71	1.42	1.12	2.01	0.61	1.16	0.59	*									
NT2RP4001739	4.83	2.71	3.87	5.22	3.09	4.63	4.39	5.41	4.57										
NT2RP4001741	10.82	7.34	4.37	12.44	9.41	10.54	7.99	6.39	5.79										
NT2RP4001753	11.73	4.55	5.91	14.42	16.38	12.01	9.64	6.92	8.76										
NT2RP4001760	12.48	7.93	6.76	6.21	7.16	7.70	2.14	2.05	1.86										
NT2RP4001787	45.15	35.87	34.25	46.58	51.77	52.14	19.69	24.09	19.41	*									
NT2RP4001790	6.06	3.59	2.88	5.91	6.74	7.08	5.5	5.13	5.27										
NT2RP4001795	25.43	15.84	22.47	18.33	17.56	16.99	11.05	11.79	10.2										
NT2RP4001803	3.51	2.55	1.55	5.77	4.36	4.15	4.43	4.25	2.97	*									
NT2RP4001805	4.04	2.46	2.43	5.53	4.54	4.71	3.91	2.99	3.66	*									
NT2RP4001809	14.99	9.07	7.27	11.92	10.72	9.25	11.36	11.25	11.16										
NT2RP4001817	16.10	8.59	7.80	8.81	9.92	9.75	5.74	6.19	5.7										
NT2RP4001822	9.90	6.09	4.79	7.82	4.55	6.51	6.73	5.44	6.61										
NT2RP4001823	1.63	1.96	0.82	1.62	2.17	1.74	1.56	1.67	0.88										
NT2RP4001827	5.09	4.68	4.45	4.54	5.32	5.79	7.53	6.64	8.76	*									
NT2RP4001828	17.04	10.89	10.46	15.89	15.47	13.14	13.38	12.00	9.76										
NT2RP4001836	5.07	3.08	3.80	4.72	5.04	5.75	5.07	4.56	2.8										
NT2RP4001838	6.83	3.89	5.07	5.21	5.01	6.41	4.27	6.56	2.85										

120/180

【0761】

【表290】

1717

1718

NT2RP4001841	5.15	2.19	2.44	6.33	5.75	3.95	4.94	4.03	3.03										
NT2RP4001842	4.08	2.37	1.90	1.96	2.08	2.74	2.12	3.59	2.22										
NT2RP4001861	19.55	11.05	8.48	18.06	19.21	17.61	12.49	10.31	10.34										
NT2RP4001877	18.38	12.98	11.71	13.65	17.92	15.26	10.17	11.03	9.86										
NT2RP4001879	6.00	4.86	5.20	4.62	6.88	7.55	4.96	6.52	5.75										
NT2RP4001889	3.83	2.48	2.26	4.36	5.15	5.12	3.39	5.09	3.84	*	+								
NT2RP4001893	4.85	2.58	3.31	5.78	4.46	6.55	5.02	4.75	1.96										
NT2RP4001896	4.86	2.86	3.13	4.46	5.44	4.95	3.44	3.93	1.91										
NT2RP4001898	12.63	7.18	6.38	11.85	13.48	14.72	8.27	7.05	8.92										
NT2RP4001901	9.37	4.10	4.58	7.22	7.41	7.58	5.92	5.84	4.25										
NT2RP4001910	44.22	14.42	25.27	36.18	28.56	31.03	15.44	16.11	13.43										
NT2RP4001925	6.01	3.53	4.07	7.13	8.88	6.52	5.38	5.68	3.89	*	+								
NT2RP4001926	5.02	2.32	4.10	6.70	3.01	7.01	3.35	4.83	1.34										
NT2RP4001927	7.81	3.22	8.37	2.90	3.77	4.75	2.11	3.46	2.61										
NT2RP4001931	12.13	7.10	9.23	9.30	11.80	10.57	7.09	9.58	5.89										
NT2RP4001933	7.27	5.93	8.24	33.37	26.48	21.53	12.07	15.48	9.59	**	+	*	+						
NT2RP4001938	11.79	6.36	5.51	7.00	8.59	7.23	7.68	7.54	9.66										
NT2RP4001942	19.13	10.55	10.00	11.76	13.07	12.47	8.35	7.90	8.71										
NT2RP4001945	3.39	2.16	1.75	1.10	2.83	1.75	3.88	3.65	3.03										
NT2RP4001946	2.78	2.76	2.10	6.68	5.62	8.03	3.2	4.28	3.28	**	+								
NT2RP4001947	0.70	0.50	0.71	3.55	3.12	4.05	1.69	2.42	0.29	**	+								
NT2RP4001950	52.07	29.14	30.34	3.90	3.31	3.63	2.85	3.53	3.23	*	-	*	-						
NT2RP4001953	6.50	3.60	5.67	12.09	12.07	9.95	5.86	6.12	3.31	**	+								
NT2RP4001966	3.87	2.06	1.81	2.93	2.33	3.06	2.56	3.55	1.61										
NT2RP4001970	18.77	7.73	6.33	7.39	9.12	8.12	6.83	7.05	6.87										
NT2RP4001975	16.12	8.35	8.50	16.73	14.38	16.13	21.64	17.08	14.87										
NT2RP4001988	6.11	2.52	2.36	2.17	2.97	2.42	4.05	5.29	6.8										
NT2RP4001996	8.88	6.41	7.06	5.35	6.06	5.33	4.86	5.72	5.5										
NT2RP4002014	5.46	3.70	3.51	5.82	4.28	3.92	5.71	6.94	6.45			*	+						
NT2RP4002018	4.51	3.12	2.83	6.79	4.88	5.98	4.66	10.23	5.14	*	+								
NT2RP4002035	6.12	4.46	6.67	7.19	6.57	6.76	5.8	5.73	6.32										
NT2RP4002043	17.40	10.99	15.66	15.62	10.19	12.89	8.93	9.28	8.15			*	-						
NT2RP4002046	6.17	4.77	3.90	3.50	9.38	4.20	6.26	5.07	7.72										
NT2RP4002047	14.83	7.78	9.72	12.74	11.68	9.86	4.4	4.88	5.22			*	-						
NT2RP4002052	3.82	2.22	2.36	3.72	2.89	4.12	4.34	4.20	5.03			*	+						
NT2RP4002056	55.72	38.98	47.46	51.12	52.01	41.19	44.9	38.97	37.38										
NT2RP4002057	17.74	8.34	10.35	10.25	6.84	10.23	9.46	9.43	9										
NT2RP4002058	5.05	3.72	3.60	3.34	2.84	3.35	3.74	3.86	2.96										
NT2RP4002064	2.43	1.64	1.15	2.53	2.72	2.44	2.13	3.96	2.74										
NT2RP4002071	6.91	5.83	6.59	9.94	11.45	10.50	6.83	7.79	5.44	**	+								
NT2RP4002075	5.65	2.21	2.77	1.76	1.64	2.01	1.03	0.80	1.27										
NT2RP4002078	12.20	5.57	6.28	21.16	11.84	9.58	9.34	5.65	7.44										
NT2RP4002081	8.20	4.41	4.38	8.71	5.52	5.98	8.56	5.96	6.86										
NT2RP4002083	1.41	0.64	0.77	1.12	0.92	0.88	1.16	1.92	2.62										
NT2RP4002099	3.50	1.74	2.24	2.98	2.94	2.77	2.69	3.45	2.97										
NT2RP4002106	16.08	11.97	16.65	14.50	11.42	13.37	8.7	8.53	7.16			*	-						
NT2RP4002111	14.95	7.66	10.77	13.64	14.70	14.45	16.75	17.55	15.81										
NT2RP4002113	5.99	2.81	3.54	4.57	5.85	6.57	6.12	5.13	4.94										
NT2RP4002116	14.14	7.04	5.48	14.30	12.58	11.93	8.14	4.91	6.09										
NT2RP4002123	15.83	9.46	8.25	6.72	6.57	5.28	1.63	2.27	1.64			*	-						
NT2RP4002126	7.11	2.89	3.58	2.17	3.83	2.41	4.77	4.31	5.35										
NT2RP4002133	10.15	4.28	5.52	6.19	8.16	5.73	5.02	6.44	4.79										
NT2RP4002136	13.83	8.55	8.39	3.76	6.14	5.01	5.63	5.28	4.82			*	-						
NT2RP4002139	23.38	27.01	30.04	25.35	29.41	24.98	13.23	24.52	19.91										
NT2RP4002174	3.31	8.15	2.46	3.71	3.16	4.15	3.14	5.41	4.47										
NT2RP4002185	10.77	7.55	7.67	15.20	13.52	13.41	10.77	8.24	8.8	*	+								

121/180

【0762】

【表291】

1719

1720

NT2RP4002186	24.35	16.62	12.92	73.40	68.88	51.66	18.99	20.53	42.77	**	+		
NT2RP4002187	16.88	9.15	8.08	13.25	18.23	17.99	14.62	13.98	23.37				
NT2RP4002189	9.49	5.18	4.64	14.52	14.99	9.78	4.92	6.78	9.43	*	+		
NT2RP4002199	3.33	0.85	1.71	2.01	2.76	1.40	1.46	4.34	2.92				
NT2RP4002206	7.79	3.61	3.56	5.56	5.23	3.75	3.53	5.24	4.66				
NT2RP4002210	3.95	1.94	2.05	3.42	2.86	2.32	2.13	4.76	2.28				
NT2RP4002222	4.87	2.50	3.89	4.48	5.59	3.24	4.1	4.89	3.82				
NT2RP4002241	10.39	8.75	9.34	8.11	10.75	7.80	3.57	5.39	6.12	**	-		
NT2RP4002248	5.75	3.15	2.68	4.58	3.49	3.31	6.08	4.55	3.57				
NT2RP4002250	2.77	1.28	0.36	1.28	1.49	1.07	2.02	0.58	1.13				
NT2RP4002259	11.44	4.70	6.93	10.37	10.26	7.96	6.18	7.00	6.72				
NT2RP4002268	9.49	7.15	6.70	7.16	8.97	8.79	12.35	10.44	12.35	*	+		
NT2RP4002288	23.22	15.06	19.08	20.88	28.68	23.53	20.52	17.25	20.1				
NT2RP4002290	9.48	5.25	5.05	15.46	15.55	18.46	13.55	11.18	12.37	**	+	*	+
NT2RP4002298	5.94	3.63	4.51	10.11	6.35	12.09	3.11	5.17	4.75				
NT2RP4002306	5.29	2.43	3.39	8.59	7.82	9.25	3.86	4.05	3.61	**	+		
NT2RP4002308	2.50	1.35	1.43	1.70	2.93	1.47	2.72	1.97	2.14				
NT2RP4002336	9.03	4.10	4.50	6.72	4.54	7.26	5.89	4.31	4.91				
NT2RP4002340	0.95	0.34	0.60	0.63	0.88	0.24	1.51	1.53	0.76				
NT2RP4002361	3.28	2.38	1.78	3.90	2.34	2.47	2.23	2.16	1.92				
NT2RP4002367	3.30	2.19	1.54	3.77	4.95	3.32	2.84	2.25	3				
NT2RP4002368	4.21	2.40	3.66	5.83	4.14	3.92	5.91	4.62	3.42				
NT2RP4002377	3.62	4.26	2.84	5.85	2.38	5.20	4.75	3.54	3.33				
NT2RP4002408	29.46	20.49	24.43	3.81	2.37	2.48	1.32	0.66	1.06	**	-	**	-
NT2RP4002425	1.74	1.67	0.75	1.77	1.60	1.39	2.92	1.48	1.25				
NT2RP4002432	8.35	5.60	3.82	5.76	5.85	4.41	8.08	6.14	6.6				
NT2RP4002447	9.10	3.90	3.22	12.78	11.88	10.40	5.91	5.47	6.48	*	+		
NT2RP4002451	2.21	2.30	1.71	3.91	4.29	3.31	5.98	6.15	6.01	**	+	**	+
NT2RP4002461	7.09	5.26	5.72	12.99	9.75	9.13	7.77	8.39	7.06	*	+		
NT2RP4002486	5.84	4.56	5.50	5.14	5.35	4.72	7.44	6.30	6.54		*	+	
NT2RP4002517	3.21	2.30	2.48	3.27	2.89	3.72	3.06	3.47	2.44				
NT2RP4002556	10.73	5.00	5.11	11.36	8.97	7.80	4.9	4.20	4.51				
NT2RP4002569	5.60	3.78	2.56	4.11	4.44	3.67	5.29	5.70	3.72				
NT2RP4002587	2.41	1.81	1.87	2.59	3.67	3.36	7.6	6.60	7.95	*	+	**	+
NT2RP4002591	7.42	6.05	5.29	12.68	12.07	10.38	7.78	4.95	7.6	**	+		
NT2RP4002607	6.11	2.67	2.59	6.08	4.47	5.73	3.49	4.31	2.91				
NT2RP4002627	5.30	4.31	4.08	5.45	8.00	6.98	9.15	7.80	7.44		**	+	
NT2RP4002638	13.62	7.50	7.90	12.59	11.82	9.24	5.81	7.23	4.46				
NT2RP4002638	3.81	2.47	2.90	6.00	2.15	4.82	6.13	6.96	4.18		*	+	
NT2RP4002639	4.77	2.18	3.85	2.27	2.26	2.48	1.79	3.34	1.18				
NT2RP4002641	8.72	3.54	3.33	4.53	5.23	4.41	5.45	5.35	8.22				
NT2RP4002658	39.52	16.53	21.90	10.69	10.22	8.63	12.92	12.25	13.09				
NT2RP4002669	8.68	5.48	3.61	6.49	4.90	5.66	4.3	4.66	5.21				
NT2RP4002677	11.90	7.10	10.78	11.62	13.84	10.32	4.5	5.32	4.63		*	-	
NT2RP4002715	6.49	4.85	5.45	16.06	11.33	12.78	11.52	13.89	13.89	**	+	**	+
NT2RP4002759	11.19	4.76	5.94	4.82	3.94	4.94	3.86	3.11	3.58				
NT2RP4002784	5.22	3.74	4.33	6.90	5.66	7.76	7.61	4.05	3.39	*	+		
NT2RP4002791	2.52	2.02	2.01	4.89	4.01	4.73	3.62	3.85	2.33	**	+		
NT2RP4002811	6.07	3.91	2.96	1.95	3.45	3.30	4.41	5.24	4.51				
NT2RP4002839	11.00	4.98	5.60	10.88	8.44	6.08	6.46	6.96	4.35				
NT2RP4002832	2.65	2.09	2.28	3.38	2.40	2.76	1.95	3.23	1.27				
NT2RP4002859	10.22	7.64	6.28	14.24	10.59	11.13	9.28	9.04	5.75				
NT2RP4002874	3.50	2.69	1.87	3.65	3.03	4.18	3.78	5.05	2.68				
NT2RP4002884	17.66	6.25	9.46	10.83	9.85	8.92	15.08	14.77	10.31				
NT2RP4002888	20.83	12.71	14.10	15.29	12.54	11.78	19.91	18.79	15.14				
NT2RP4002891	6.49	3.33	5.04	17.64	15.92	12.46	8.11	7.56	7.35	**	+	*	+

122/180

1721

1722

NT2RP4002894	30.47	15.42	16.30	15.33	13.44	13.63	14.61	7.84	11.34										
NT2RP4002896	5.01	2.57	1.03	5.77	4.90	3.35	4.85	5.20	6.5										
NT2RP4002905	3.65	2.18	2.47	3.73	2.63	3.46	2.67	3.64	2.22										
NT2RP4002907	6.79	1.23	2.84	16.01	14.42	10.02	12.06	10.10	6.54	*	+								
NT2RP5003459	65.35	36.44	48.17	27.67	30.09	25.05	9.64	20.91	22.09										
NT2RP5003461	4.58	3.60	3.17	6.87	4.80	7.46	3.05	4.17	2.86	*	+								
NT2RP5003471	5.96	3.26	3.68	5.59	5.78	6.38	36.49	36.45	36.48										
NT2RP5003477	4.19	2.26	3.16	4.58	5.06	6.58	6.46	4.38	3.4										
NT2RP5003487	220.55	93.22	98.28	181.60	187.80	154.38	86.45	85.87	93.23										
NT2RP5003492	7.41	4.46	3.61	6.80	6.09	7.24	6.01	5.67	4.83										
NT2RP5003500	3.73	2.01	1.80	4.33	3.62	5.68	2.91	3.93	3.14										
NT2RP5003506	9.63	4.24	5.17	6.58	8.38	7.49	5.4	7.54	7.66										
NT2RP5003512	2.05	1.82	0.90	1.93	2.76	1.89	1.76	3.04	2.68										
NT2RP5003522	5.00	3.31	4.09	6.05	5.02	4.70	4.69	4.96	3.11										
NT2RP5003524	2.66	1.03	1.85	3.05	3.14	2.14	2.01	1.80	0.86										
NT2RP5003527	27.32	17.39	20.11	33.15	29.19	33.15	34.18	28.33	30.99	*	+								
NT2RP5003531	6.09	4.05	3.52	14.63	15.87	11.17	18.91	10.15	13.33	**	+								
NT2RP5003534	4.69	3.24	2.48	4.56	5.46	3.21	3.85	3.74	4.1										
NT2RP6000020	14.93	5.50	7.94	19.43	12.24	14.47	28	17.69	22.01										
NT2RP6000023	2.09	1.92	1.10	2.89	3.69	3.48	1.85	3.95	3.04	*	+								
NT2RP6000020	6.72	2.85	2.69	5.15	4.13	6.91	3.13	4.74	4.15										
NT2RP6000063	4.32	1.86	2.74	4.12	3.95	5.49	4.77	3.84	5.17										
NT2RP6000074	7.65	3.63	3.82	5.82	4.62	5.47	3.91	5.25	4.12										
NT2RP6000083	7.65	4.46	4.22	5.62	7.05	9.12	4.96	6.80	6.49										
NT2RP6000100	8.20	3.69	3.69	11.31	10.03	10.20	5.69	6.11	4.22	*	+								
NT2RP6000123	8.42	4.03	3.87	7.40	6.54	4.76	5.08	5.14	4.33										
NT2RP6000129	5.14	2.45	3.11	3.95	4.30	4.21	3.96	4.16	4.57										
NT2RP6000147	3.79	2.50	3.26	15.24	15.27	11.86	26.48	14.22	25.1	**	+								
NT2RP6000163	1.43	1.14	1.15	3.25	1.30	2.00	1.02	2.54	1.73										
NT2RP6000181	7.19	4.67	4.25	6.16	6.80	4.73	6.67	3.10	6.2										
NT2RP6000182	5.25	3.12	3.43	5.76	4.23	7.79	3.45	3.70	2.44										
QVARC1000001	4.47	2.05	2.92	5.01	4.27	3.71	5.92	4.78	4.37										
QVARC1000003	4.03	2.27	2.17	3.53	4.26	1.98	1.87	2.81	4.16										
QVARC1000004	69.94	45.81	40.28	31.28	33.52	34.13	14.2	20.99	22.91	*	+								
QVARC1000006	2.75	1.60	1.91	3.55	3.17	2.27	3.59	3.71	3.52										
QVARC1000013	3.58	2.31	1.87	3.88	4.15	3.20	3.52	4.55	2.95										
QVARC1000014	5.72	2.95	3.69	6.24	6.32	5.61	4.07	4.99	4.34										
QVARC1000017	6.14	3.05	3.33	4.90	5.12	5.05	3.15	5.17	5.31										
QVARC1000026	55.69	36.49	45.68	51.02	60.13	48.46	28.42	36.95	25.22										
QVARC1000035	9.77	8.46	8.93	13.12	14.00	9.30	7.02	5.89	5.3										
QVARC1000037	31.27	16.99	12.47	49.92	29.93	32.59	18.22	25.08	32.08										
QVARC1000038	10.77	5.52	3.11	12.87	13.32	13.63	6.74	5.82	8.66	*	+								
QVARC1000060	3.24	1.54	1.26	3.04	2.70	2.45	2.09	2.66	3.05										
QVARC1000068	2.38	1.15	1.10	3.07	2.77	1.87	1.01	3.23	1.66										
QVARC1000069	4.64	2.24	2.58	7.95	8.04	5.29	4.94	7.33	5.21	*	+								
QVARC1000071	4.18	2.24	2.19	3.21	4.19	2.93	1.32	4.38	1.25										
QVARC1000073	116.66	59.06	70.03	104.67	109.44	102.05	127.1	180.67	194.9										
QVARC1000083	16.13	9.03	10.85	16.27	15.52	17.85	9.32	13.62	11.15										
QVARC1000085	90.31	52.35	57.44	84.93	91.25	74.75	46.89	55.51	55.51										
QVARC1000086	3.63	2.07	4.18	7.09	7.77	8.13	5.87	6.77	6.77	**	+								
QVARC1000087	2.46	0.70	0.93	1.65	1.80	2.44	2.22	3.58	3.58										
QVARC1000090	7.22	4.69	6.24	15.64	14.18	15.90	5.67	9.11	9.11	**	+								
QVARC1000091	3.66	1.42	2.09	5.7	4.57	5.66	4.01	3.77	3.77	*	+								
QVARC1000092	3.91	1.98	2.18	6.09	6.36	8.26	4.35	4.86	4.86	*	+								
QVARC1000105	11.95	8.25	9.35	12.3	11.58	13.87	6.66	8.05	8.05										
QVARC1000106	23.29	10.32	10.91	20.75	17.39	12.69	12.13	18.29	18.29										

123/180

【0764】

【表293】

1724

OVARC1000109	10.73	4.48	6.00	9.44	8.48	8.37	6.70	8.07	8.07				
OVARC1000113	4.43	3.28	2.32	5.28	7.68	6.28	3.04	3.01	3.01	*	+		
OVARC1000114	4.61	1.83	2.98	6.68	7.52	8.77	4.82	5.56	5.56	*	+		
OVARC1000133	2.28	0.62	2.11	1.97	3.23	1.32	1.31	3.42	3.42				
OVARC1000137	7.57	3.31	3.78	7.45	5.43	6.40	5.03	9.51	9.51				
OVARC1000139	8.5	5.04	5.90	7.42	5.19	7.20	5.43	7.04	7.04				
OVARC1000145	1.66	0.51	1.26	2.03	2.15	2.60	1.95	1.95	1.95	*	+		
OVARC1000148	13.99	5.79	5.64	16.54	19.40	9.14	7.33	8.83	8.83				
OVARC1000151	5.62	2.25	3.47	4.79	5.94	4.15	4.17	6.14	6.14				
OVARC1000157	5.78	3.92	3.63	20.18	23.53	19.12	7.05	10.69	10.69	**	+	*	+
OVARC1000162	1.04	0.27	1.30	1.82	2.05	0.82	1.71	1.67	1.67				
OVARC1000168	6.93	3.43	5.38	9.14	7.70	8.50	5.44	8.50	8.50	*	+		
OVARC1000169	20.78	9.01	10.52	18.85	14.31	18.81	15.67	26.42	26.42				
OVARC1000178	6.27	4.19	5.21	6.05	5.93	6.06	4.30	3.93	3.93				
OVARC1000182	1.08	0.33	0.60	3.18	1.53	2.07	1.58	1.16	1.16	*	+		
OVARC1000186	11.87	6.09	4.34	4.72	8.03	4.57	4.49	8.00	8				
OVARC1000188	6.88	3.30	4.11	6.26	4.11	4.48	4.18	5.80	5.8				
OVARC1000191	2.39	0.93	1.25	1.87	4.24	1.53	1.02	3.43	3.43				
OVARC1000198	7.48	2.50	4.22	12.55	13.51	9.27	4.79	6.14	6.14	*	+		
OVARC1000208	7.66	3.85	6.83	11.11	11.76	10.78	8.71	6.63	6.63	**	+		
OVARC1000209	5.19	2.21	3.10	4.98	5.19	3.99	3.67	6.12	6.12				
OVARC1000212	7.76	3.64	5.91	6.62	4.86	7.76	4.09	6.97	6.97				
OVARC1000216	1.71	1.54	1.80	2.95	1.87	2.06	1.88	2.20	2.2			*	+
OVARC1000240	9.19	4.82	3.93	10.89	11.55	7.32	4.66	6.08	6.08				
OVARC1000241	8.4	2.88	3.50	6.97	5.95	3.69	4.83	5.66	5.66				
OVARC1000249	5.89	2.71	3.55	5.91	2.56	3.50	4.13	5.08	5.08				
OVARC1000284	16.05	11.01	13.12	50.15	29.76	29.83	42.38	33.82	33.82	*	++		
OVARC1000255	5.5	3.14	2.99	5.45	4.17	3.19	3.91	4.30	4.3				
OVARC1000267	8.95	5.90	5.53	9.61	7.91	10.70	8.96	10.59	10.59				
OVARC1000275	0.38	0.28	0.65	1.7	1.69	1.90	1.31	9.09	9.09	**	+	**	+
OVARC1000287	2.16	1.07	1.61	5.38	6.97	4.90	26.09	33.14	33.14	**	+	**	+
OVARC1000288	7.19	3.43	4.43	6.36	6.18	3.91	4.34	4.81	4.81				
OVARC1000290	8.86	6.47	4.36	11.32	12.55	7.25	6.14	7.12	7.12				
OVARC1000382	3.96	1.75	1.50	3.75	4.71	3.28	2.04	3.19	3.19				
OVARC1000394	6.08	4.82	3.98	7.97	7.57	5.26	4.58	6.93	6.93				
OVARC1000397	5.1	1.95	3.30	4.25	2.68	4.18	3.69	3.54	3.54				
OVARC1000399	6.17	3.11	3.95	6.94	5.55	4.98	5.49	5.61	5.61				
OVARC1000512	4.47	2.31	2.62	3.43	3.39	3.03	5.14	4.44	4.44				
OVARC1000513	7.23	3.04	5.41	6.92	6.31	4.37	7.31	10.70	10.7				
OVARC1000521	8.81	5.88	6.66	13.97	15.87	13.56	14.26	12.53	12.53	**	+	**	+
OVARC1000526	3.94	3.57	2.28	3.59	3.18	3.94	3.62	3.71	3.71				
OVARC1000527	4.66	2.13	3.59	7.38	4.82	4.34	3.97	5.68	5.68				
OVARC1000531	6.82	4.80	4.04	7.15	6.72	8.39	4.61	6.40	6.4				
OVARC1000535	3.22	3.45	3.68	6.19	5.78	6.01	4.99	5.32	5.32	*	+		
OVARC1000547	2.86	2.21	1.39	1.74	2.06	3.33	1.79	3.03	3.03				
OVARC1000548	7.01	4.29	4.68	13.43	12.42	16.47	7.65	8.17	8.17	**	+	*	+
OVARC1000563	4.22	3.97	3.08	6.15	6.28	7.74	2.83	4.38	4.38	**	+		
OVARC1000577	2.82	1.76	1.53	3.08	2.53	1.71	0.35	2.23	2.23				
OVARC1000842	5.76	1.98	3.91	4.79	5.06	3.60	4.10	6.90	6.9				
OVARC1000864	6.02	5.30	4.11	6.76	8.20	10.33	8.85	9.44	9.44	*	++		
OVARC1000901	2.8	1.75	1.96	2.86	2.28	2.89	2.67	3.48	3.48				
OVARC1000906	114.78	80.20	88.37	88.01	73.54	119.34	90.62	95.77	95.77				
OVARC1000907	4.6	3.44	3.17	5.58	5.61	8.84	5.08	4.38	4.38				
OVARC1000908	16.3	13.53	12.64	45.51	42.22	49.78	26.85	32.12	32.12	**	+	**	+
OVARC10009410	6.71	4.55	5.34	4.6	7.22	6.74	6.55	6.11	6.11				
OVARC10009411	3.32	1.84	2.60	3.91	7.15	2.94	1.78	2.39	2.39				

1725

1726

OVARC100414	2.94	2.41	3.01	5.83	4.82	5.60	3.16	3.78	3.78	**	+	*	+
OVARC100420	11.4	6.17	7.59	9.91	9.38	10.06	10.09	13.16	13.16				
OVARC100431	8.6	6.78	5.53	8.33	7.86	10.75	8.17	6.59	6.59				
OVARC100437	3.68	2.71	4.36	3.26	4.27	4.49	3.23	3.96	3.96				
OVARC100431	28.24	22.85	26.14	17.5	18.78	21.85	14.12	15.50	15.5	*	-	**	-
OVARC100437	4.74	2.97	4.16	5.12	6.15	7.20	4.22	6.60	6.6				
OVARC100439	7.31	6.90	5.38	7.44	6.69	8.00	6.48	4.83	4.83				
OVARC100440	10.79	6.84	6.93	7.88	7.24	7.80	6.48	7.22	7.22				
OVARC100442	5.47	3.48	2.90	10.37	8.05	7.61	4.21	5.71	5.71	*	+		
OVARC100443	2.37	1.87	2.77	3.52	3.55	4.55	2.82	6.19	6.19	*	+		
OVARC100441	3.39	2.34	2.79	3.41	2.83	2.56	4.13	3.34	3.34				
OVARC100445	4.49	3.75	4.70	4.63	4.57	4.49	3.93	2.86	2.86				
OVARC100446	5.63	3.82	4.46	5.01	4.97	7.62	6.00	5.12	5.12				
OVARC100447	3.64	2.33	2.91	3.88	3.66	4.53	4.40	4.32	4.32			*	+
OVARC100470	4.4	2.42	1.89	7.76	7.31	7.37	4.36	3.86	3.86	**	+		
OVARC100473	5.77	6.12	2.59	5.13	4.08	6.65	4.72	6.17	6.17				
OVARC100479	10.65	6.40	6.55	8.36	8.23	12.25	7.74	6.99	6.99				
OVARC100484	7.73	3.54	4.68	14.41	17.12	13.60	11.04	9.93	9.93	**	+	*	+
OVARC100486	3.13	1.48	1.74	5.56	5.39	7.63	4.10	3.04	3.04	**	+		
OVARC100486	0.32	0.95	1.13	0.23	0.59	1.74	1.38	0.85	0.85				
OVARC100520	0.79	1.22	1.43	1.76	1.97	2.08	2.17	1.68	1.68	*	+	*	+
OVARC100522	4.89	4.05	3.21	7.99	8.63	12.13	8.58	8.73	8.73	*	**	+	+
OVARC100526	5.23	3.76	3.40	9.44	8.41	9.79	6.60	6.83	6.83	**	+	**	+
OVARC100529	8.29	3.03	3.79	8.43	8.08	7.91	6.33	6.00	6				
OVARC100533	13.85	10.76	9.50	10.46	10.65	9.69	10.80	10.74	10.74				
OVARC100543	2.14	1.23	0.78	1.99	1.06	1.67	1.34	1.95	1.95				
OVARC100550	3.91	2.99	2.96	3.41	5.27	5.08	3.89	3.69	3.69				
OVARC100553	7.96	6.39	6.63	10.34	11.92	12.52	8.20	8.94	8.94	**	+	*	+
OVARC100556	2.91	2.73	2.33	4.64	4.36	4.57	3.30	5.76	5.76	**	+		
OVARC100557	1.8	2.00	2.08	3.66	2.89	3.58	2.75	2.23	2.23	**	+		
OVARC100561	5.49	5.12	4.27	12.79	11.21	12.35	4.34	6.50	6.5	**	+		
OVARC100564	11	4.97	4.69	6.39	9.03	5.47	5.12	5.72	5.72				
OVARC100573	3.43	1.54	1.73	4.84	5.71	5.20	3.22	2.70	2.7	**	+		
OVARC100576	22.35	9.42	12.58	14.84	14.82	13.96	18.96	21.39	21.39				
OVARC100578	3.78	1.92	1.91	7.25	4.00	7.95	3.26	3.45	3.45	*	+		
OVARC100581	2.32	0.98	1.31	2.39	2.02	2.50	0.87	2.36	2.36				
OVARC100586	4.15	3.94	3.82	5.69	4.46	5.03	7.98	9.37	9.37	*	**	+	+
OVARC100588	3.09	2.32	2.34	6.24	5.07	6.64	3.10	4.00	4	**	+	*	+
OVARC100605	3.48	1.27	1.57	3.94	3.34	1.96	2.17	3.54	3.54				
OVARC100622	16.94	7.82	7.29	28.21	27.34	23.72	13.10	15.48	15.48	*	+		
OVARC100636	7.07	3.14	2.94	8.06	7.46	5.78	4.15	5.40	5.4				
OVARC100640	1.93	1.10	2.17	2.95	3.95	2.11	1.86	2.87	2.87				
OVARC100642	6.55	3.47	4.45	4.81	4.28	3.96	4.42	5.10	5.1				
OVARC100641	8.83	4.09	5.47	7.42	7.41	7.08	6.76	7.35	7.35				
OVARC100677	5.49	3.25	4.84	5.23	4.19	5.75	3.47	4.95	4.95				
OVARC100678	3.24	2.45	2.41	6.56	3.68	3.55	3.07	3.23	3.23				
OVARC100679	2.29	2.05	2.51	5.5	7.69	3.61	2.67	3.02	3.02	*	+	*	+
OVARC100681	3.04	1.58	2.55	2.83	3.65	2.16	1.27	3.32	3.32				
OVARC100682	5.34	2.89	3.15	10.89	12.80	6.98	6.81	6.70	6.7	*	+	*	+
OVARC100689	6.35	2.64	5.24	6.82	5.05	3.15	3.89	5.86	5.86				
OVARC100700	4.87	2.36	3.84	6.43	6.33	5.87	5.58	3.71	3.71	*	+		
OVARC100703	6.09	5.10	4.50	10.85	8.68	9.16	5.32	6.24	6.24	**	+		
OVARC100722	6.99	3.21	3.22	7.7	5.28	6.60	3.85	5.25	5.25				
OVARC100724	12.55	5.82	7.48	9.62	7.07	8.57	9.99	8.90	8.9				
OVARC100727	8.32	3.91	3.99	6.93	6.40	4.72	3.99	5.01	5.01				
OVARC100730	6.1	3.39	3.84	6.3	8.93	6.59	2.98	3.46	3.46				

125/180

【0766】

【表295】

1727

1728

QVARC1000741	7.47	3.93	4.05	6.71	8.34	4.11	5.58	6.82	6.82				
QVARC1000744	2.7	1.49	1.95	3.42	4.80	3.86	2.21	3.03	3.03	*			
QVARC1000764	9.15	7.18	6.72	6.1	5.81	7.27	6.23	6.93	6.93				
QVARC1000769	1.96	2.22	1.65	4.18	3.56	4.40	2.93	2.93	2.93	**	+	**	+
QVARC1000771	3.36	1.52	2.49	4.38	3.35	3.58	3.00	4.34	4.34				
QVARC1000773	223.93	75.55	197.24	131.33	115.24	132.74	69.02	82.73	82.73				
QVARC1000775	5.89	2.38	2.57	10.9	11.89	6.67	5.95	7.36	7.36	*	+		
QVARC1000778	5.16	2.89	2.70	7.19	7.19	4.94	4.21	3.79	3.79				
QVARC1000779	1.34	0.25	1.68	0.81	2.17	1.66	0.98	2.78	2.78				
QVARC1000781	3.01	1.11	1.81	3.21	4.12	3.42	2.43	3.96	3.96				
QVARC1000787	5.12	1.26	2.40	6.21	4.91	6.16	2.68	3.80	3.8				
QVARC1000789	17.92	12.51	11.26	12.68	11.30	14.18	7.52	8.71	8.71				
QVARC1000809	10.27	6.21	6.25	13.32	11.12	11.87	8.07	9.42	9.42	*	+		
QVARC1000802	3.94	1.53	1.34	4.85	5.51	3.97	3.28	3.23	3.23				
QVARC1000819	7.31	2.74	2.89	9.23	8.19	6.66	4.42	6.46	6.46				
QVARC1000811	4.94	1.49	1.98	3.69	5.14	3.20	2.80	3.11	3.11				
QVARC1000814	8.98	4.85	4.30	12.34	14.84	13.49	5.29	9.28	9.28	*	+		
QVARC1000816	5.53	2.23	3.34	6.25	6.38	4.13	4.96	10.86	10.86				
QVARC1000817	0.67	0.84	0.17	1.03	1.43	0.88	1.03	1.18	1.18				
QVARC1000834	7.9	3.52	4.48	7.01	4.99	6.90	5.30	8.11	8.11				
QVARC1000846	8.76	5.89	5.62	13.13	13.07	12.45	7.92	8.86	8.86	**	+		
QVARC1000850	4.53	4.35	3.79	5.06	4.86	6.51	5.09	5.93	5.93		*	+	
QVARC1000853	10.26	6.75	7.96	17.45	22.42	13.77	15.27	17.34	17.34	*	+	**	+
QVARC1000863	2.31	1.51	1.67	2.98	3.34	3.48	2.84	3.92	3.92	**	+	*	+
QVARC1000873	5.08	3.94	3.56	7.67	7.81	9.71	8.49	9.22	9.22	**	+	**	+
QVARC1000875	13.15	7.32	6.94	10.33	8.49	11.65	7.63	12.92	12.92	**	+	**	+
QVARC1000876	3.56	1.95	2.71	3.83	2.75	3.80	2.91	3.90	3.9				
QVARC1000883	11.24	5.79	7.03	7.42	6.63	8.18	6.12	10.30	10.3				
QVARC1000885	1.99	1.85	0.96	2.91	2.72	4.05	1.81	1.84	1.84	*	+		
QVARC1000886	3.79	3.90	3.30	5.23	4.59	3.88	4.18	4.19	4.19		*	+	
QVARC1000890	16.12	9.23	8.22	13.23	13.98	12.06	8.12	8.78	8.78				
QVARC1000891	9.14	4.38	8.52	6.77	7.63	5.67	3.14	5.82	5.82				
QVARC1000897	1.42	0.51	0.89	0.57	0.73	1.37	0.82	2.14	2.14				
QVARC1000912	3.17	1.30	1.93	1.64	2.12	2.69	2.76	3.24	3.24				
QVARC1000914	1.78	1.84	1.59	1.53	2.20	2.36	1.62	3.25	3.25				
QVARC1000915	6.15	3.81	2.82	7.18	8.08	11.76	6.61	6.54	6.54				
QVARC1000916	3.99	3.47	3.85	4.78	5.13	5.34	5.36	5.24	5.24	**	+	**	+
QVARC1000924	3.43	2.14	2.20	3.95	5.25	6.94	3.56	2.87	2.87	*	+		
QVARC1000928	2.3	1.45	1.90	2.94	5.21	2.82	3.59	4.30	4.3		**	+	
QVARC1000936	2.24	1.71	1.41	6.64	6.60	5.33	2.20	4.51	4.51	**	+		
QVARC1000937	4.37	3.36	3.65	4.37	4.24	5.18	4.25	6.07	6.07				
QVARC1000945	6.45	6.49	5.55	5.48	7.78	7.98	6.71	7.64	7.64				
QVARC1000948	0.7	1.14	1.80	0.8	1.44	1.73	1.79	1.83	1.83				
QVARC1000956	4.51	3.81	4.84	5.21	4.57	6.49	5.02	4.22	4.22				
QVARC1000959	3.91	3.29	2.51	6.03	6.37	10.71	4.34	3.63	3.63	*	+		
QVARC1000960	15.58	12.49	8.74	25.21	26.65	32.56	13.09	13.48	13.48	**	+		
QVARC1000964	7.4	4.78	4.49	6.36	5.85	5.60	6.26	4.60	4.6				
QVARC1000971	2.15	0.88	1.23	2.29	3.01	2.68	1.01	1.16	1.16	*	+		
QVARC1000975	3.59	2.39	2.50	3.12	3.77	3.56	3.45	3.78	3.78				
QVARC1000976	1.43	1.10	0.67	1.47	1.70	1.61	1.96	1.76	1.76		*	+	
QVARC1000981	4.17	3.59	2.08	3.86	4.73	5.67	6.53	5.87	5.87		*	+	
QVARC1000982	5.28	3.08	4.53	3.99	3.64	7.26	3.94	5.23	5.23				
QVARC1000984	2.89	2.32	3.02	4.83	4.31	6.06	3.98	4.72	4.72	*	+	**	+
QVARC1000985	6.28	3.58	3.62	10.03	11.01	12.54	6.34	8.81	8.81	**	+	*	+
QVARC1000986	2.44	0.93	1.29	3.23	3.01	2.60	1.59	2.15	2.15	*	+		
QVARC1000999	13.81	7.04	7.16	18.71	16.83	17.82	9.89	10.20	10.2	*	+		

126/180

【0767】

【表296】

1729

1730

OVARC1001000	10.01	7.69	7.61	19.45	23.56	18.96	10.07	11.74	11.74	**	+	*	+
OVARC1001004	1.03	0.80	0.91	1.57	2.14	1.61	1.90	1.48	1.48	*	+	*	+
OVARC1001010	1.8	1.08	0.56	1.62	1.36	2.03	1.35	1.40	1.4				
OVARC1001011	3.43	2.88	3.13	3.51	3.30	4.55	2.89	3.10	3.1				
OVARC1001030	38.32	24.93	30.71	46.79	41.55	50.96	53.76	59.72	59.72	*	+	**	+
OVARC1001032	1.55	1.32	1.67	3.18	2.58	2.77	2.83	1.37	1.37	**	+		
OVARC1001034	2.4	1.70	2.13	3.14	3.10	3.44	2.01	2.66	2.66	**	+		
OVARC1001038	12.68	9.34	7.92	11.12	13.30	12.41	6.75	6.49	6.49				
OVARC1001040	8.91	6.59	4.66	14.02	14.04	19.13	7.93	7.81	7.81	*	+		
OVARC1001041	6.31	3.56	4.31	8.16	10.01	10.61	5.62	4.95	4.95	*	+		
OVARC1001044	1.81	1.80	2.22	2.71	2.48	2.79	2.22	2.94	2.94	*	+		
OVARC1001049	9.39	8.47	8.39	15.62	16.10	16.18	9.93	8.69	8.69	**	+		
OVARC1001051	57.5	54.01	57.15	51.44	56.52	72.78	36.05	33.73	33.73	**	+		
OVARC1001054	1.32	1.27	1.50	2.46	1.80	2.94	1.81	1.58	1.58	*	+	*	+
OVARC1001055	3.77	1.65	2.45	4.24	4.50	2.62	2.94	3.56	3.56				
OVARC1001062	11.74	5.75	4.85	11.81	10.68	10.78	3.12	5.25	5.25				
OVARC1001063	1.99	1.18	1.96	2.64	2.00	1.58	1.32	1.86	1.86				
OVARC1001068	6.51	2.07	3.30	4.91	4.25	4.95	3.64	6.26	6.26				
OVARC1001072	9.32	6.54	7.65	10.21	8.94	9.18	6.17	9.88	9.88				
OVARC1001073	3.46	0.94	2.36	3.97	3.24	3.42	2.17	2.06	2.06				
OVARC1001074	1.75	0.40	1.35	1.71	2.05	2.60	0.86	1.25	1.25				
OVARC1001078	7.1	3.90	5.62	11.77	8.65	7.84	4.87	6.07	6.07				
OVARC1001083	5.2	2.42	3.41	5.59	4.12	3.31	4.28	6.32	6.32				
OVARC1001086	5.76	2.45	2.47	3.85	5.26	3.78	2.15	3.47	3.47				
OVARC1001091	3.91	3.54	2.95	5.93	5.39	4.15	4.20	3.41	3.41	*	+		
OVARC1001092	4.33	2.96	3.51	6.04	6.34	5.90	3.69	5.56	5.56	**	+		
OVARC1001104	1.53	0.53	0.40	1.32	1.57	1.20	0.63	1.14	1.14				
OVARC1001107	9.82	5.46	6.15	6.8	4.45	8.10	6.28	6.79	6.79				
OVARC1001113	4.68	3.14	2.92	4.82	4.00	4.79	2.64	3.74	3.74				
OVARC1001117	6.69	2.96	3.38	8.53	8.56	12.29	4.84	6.33	6.33	*	+		
OVARC1001118	8.12	5.06	4.70	11.61	11.15	10.02	5.36	7.35	7.35	*	+		
OVARC1001125	18.96	12.37	9.61	15.08	18.61	12.67	4.50	5.26	5.26			*	-
OVARC1001129	5.21	3.98	5.45	6.68	4.55	3.29	2.17	3.47	3.47	*	+	*	-
OVARC1001132	6.52	3.70	5.55	7.12	8.81	9.06	2.18	2.72	2.72	*	+	*	-
OVARC1001138	16.11	12.56	10.50	16.95	13.15	17.48	16.11	18.55	18.55				
OVARC1001141	5.54	2.36	3.55	4.59	3.46	4.09	3.37	5.02	5.02				
OVARC1001144	5.08	2.38	3.52	7.23	5.71	6.41	6.14	7.71	7.71	*	+	*	+
OVARC1001161	5.7	2.64	4.14	8.62	7.37	7.00	3.80	4.51	4.51	*	+	*	+
OVARC1001162	7.21	3.90	4.19	8.88	8.61	6.05	4.92	4.79	4.79				
OVARC1001163	8.43	4.40	4.84	6.45	6.12	5.05	5.16	8.27	8.27				
OVARC1001167	6.39	2.75	3.96	9.57	10.93	6.52	6.33	5.84	5.84				
OVARC1001169	2.12	0.82	1.00	1.91	2.68	3.48	1.25	1.14	1.14				
OVARC1001170	5.03	2.13	3.01	9.37	9.52	8.69	6.25	6.17	6.17	**	+	*	+
OVARC1001171	13.87	7.94	9.22	17.9	10.00	17.22	8.13	7.41	7.41				
OVARC1001173	6.07	4.57	5.00	13.94	11.16	14.09	6.22	8.18	8.18	**	+	*	+
OVARC1001176	120.6	80.54	85.77	70.13	72.81	62.53	40.27	47.88	47.88	*	+	*	-
OVARC1001180	11.62	7.41	6.61	16.44	18.19	11.30	11.76	9.48	9.48				
OVARC1001188	6.48	2.62	2.03	5.02	7.63	5.54	3.93	4.15	4.15				
OVARC1001200	2.22	1.30	1.20	5.74	5.77	3.98	2.73	3.72	3.72	**	+	*	+
OVARC1001203	7.54	4.54	7.94	10.59	9.48	9.04	5.59	7.44	7.44				
OVARC1001206	4.56	1.77	2.35	5.27	2.32	4.40	4.18	3.01	3.01				
OVARC1001209	5.41	4.08	4.25	4.84	4.05	4.40	5.08	5.10	5.1				
OVARC1001219	2.78	1.08	2.61	2.53	1.72	2.02	0.89	2.02	2.02				
OVARC1001222	2.69	0.99	2.05	4.53	4.36	2.81	4.34	4.56	4.56			**	+
OVARC1001232	6.79	3.23	4.22	10.1	7.69	5.83	3.19	6.06	6.06				
OVARC1001240	5.42	2.74	3.04	7.44	8.60	6.15	4.85	5.14	5.14	*	+		

127/180

1731

1732

OVARC001243	1.72	1.35	1.37	1.54	2.52	1.73	1.36	2.41	2.41								
OVARC001244	24.7	9.04	13.89	22.81	23.41	15.18	12.84	15.77	15.77								
OVARC001246	40.74	22.08	30.73	92.94	72.86	54.67	53.93	71.88	71.88	*	*	*	*	*	*	*	*
OVARC001247	8.36	4.54	5.70	8.31	7.58	6.86	6.44	6.70	6.7								
OVARC001260	5.56	1.98	3.43	3.72	4.11	5.56	3.81	5.29	5.29								
OVARC001261	7.49	5.34	5.88	8.27	8.14	6.50	4.18	3.66	3.66		*	*	*	*	*	*	*
OVARC001266	9.66	6.34	6.78	20.35	19.09	14.70	18.61	12.90	12.9	**	*	*	*	*	*	*	*
OVARC001270	2.46	0.92	1.16	1.01	0.99	1.69	1.24	1.98	1.98								
OVARC001271	7.39	3.05	5.29	8.27	10.72	9.05	7.37	6.66	6.66	*	*	*	*	*	*	*	*
OVARC001282	1.01	0.92	0.97	0.97	2.26	1.76	1.02	2.02	2.02								
OVARC001326	2.46	1.56	1.43	2.56	2.90	3.81	2.32	2.50	2.5								
OVARC001306	7.3	3.30	5.02	6.03	4.37	5.50	5.45	6.39	6.39								
OVARC001314	0.91	0.46	0.79	1.37	1.95	2.32	1.59	1.62	1.62	*	*	*	*	*	*	*	*
OVARC001316	1.39	0.64	0.79	0.83	1.74	1.83	1.60	1.04	1.04								
OVARC001329	14.48	8.75	10.66	26.47	22.48	16.87	10.91	14.31	14.31	*	*	*	*	*	*	*	*
OVARC001330	5.69	3.01	1.92	3.71	3.31	3.24	2.35	2.85	2.85								
OVARC001336	5.35	4.02	3.78	4.8	5.04	6.17	4.16	5.22	5.22								
OVARC001338	3	2.42	3.08	2.63	3.26	3.21	2.60	4.03	4.03								
OVARC001339	18.39	11.67	11.13	15.76	12.03	15.86	13.83	17.02	17.02								
OVARC001340	3.7	2.44	2.40	2.48	2.50	2.72	1.64	1.40	1.4		*	*	*	*	*	*	*
OVARC001341	9.61	7.33	5.62	10.7	12.45	13.37	7.41	10.65	10.65	*	*	*	*	*	*	*	*
OVARC001342	133.57	112.33	102.75	148.81	134.63	172.83	71.00	44.68	44.68		**	*	*	*	*	*	*
OVARC001344	7.19	4.91	4.20	12.04	11.73	10.02	5.70	6.29	6.29	**	*	*	*	*	*	*	*
OVARC001357	1.77	0.51	0.85	0.71	1.22	1.30	1.05	2.71	2.71								
OVARC001359	12.91	9.14	12.19	10.45	11.07	11.24	11.72	11.75	11.75								
OVARC001360	1.13	0.79	1.43	0.68	1.47	0.77	1.27	2.96	2.96								
OVARC001369	3.18	3.27	2.79	3.55	2.73	3.58	3.69	3.39	3.39								
OVARC001372	2.77	2.30	1.69	2.23	2.48	3.94	3.04	2.69	2.69								
OVARC001376	2.87	2.00	1.97	5.27	5.80	7.45	3.84	3.47	3.47	**	*	*	*	*	*	*	*
OVARC001381	9.02	7.72	5.78	16.38	17.31	19.84	9.24	7.41	7.41	**	*	*	*	*	*	*	*
OVARC001391	4.51	2.73	2.85	3.51	4.11	3.13	3.49	3.91	3.91								
OVARC001392	8.74	6.58	5.89	10.76	13.40	11.71	12.35	14.18	14.18	*	*	*	*	*	*	*	*
OVARC001399	8.85	5.58	4.72	7.92	8.25	8.82	4.81	5.40	5.4								
OVARC001417	2.7	1.43	2.23	1.21	1.52	2.52	2.51	2.99	2.99								
OVARC001419	4.3	5.24	4.00	3.68	3.86	6.94	5.84	6.00	6		*	*	*	*	*	*	*
OVARC001425	2.29	2.40	2.49	3.29	2.74	4.54	3.29	3.09	3.09		**	*	*	*	*	*	*
OVARC001436	2.31	2.50	1.77	3.81	3.30	4.11	3.38	2.41	2.41	**	*	*	*	*	*	*	*
OVARC001442	3.28	3.48	2.35	2.21	3.99	4.48	3.98	3.31	3.31								
OVARC001451	2.33	1.90	1.35	3.6	3.77	3.76	1.55	1.55	1.55	**	*	*	*	*	*	*	*
OVARC001462	3.08	2.65	1.79	3.37	3.43	2.89	2.90	3.86	3.86								
OVARC001463	1.36	0.57	0.90	1.69	3.97	2.45	2.96	1.73	1.73	**	*	*	*	*	*	*	*
OVARC001476	9.08	6.86	7.98	15.11	12.70	14.85	28.29	23.49	23.49	**	*	*	*	*	*	*	*
OVARC001480	2.63	2.84	2.87	3.18	2.98	4.97	4.13	4.00	4								
OVARC001489	0.44	0.69	0.81	2.69	2.29	3.27	1.10	4.03	4.03	**	*	*	*	*	*	*	*
OVARC001493	1.25	1.74	1.87	2.29	2.11	2.40	3.16	2.54	2.54	*	*	*	*	*	*	*	*
OVARC001496	8.58	6.56	5.62	10.89	7.25	13.93	7.36	6.38	6.38								
OVARC001499	2.77	1.81	1.79	9.3	11.43	8.77	4.71	4.12	4.12	**	*	*	*	*	*	*	*
OVARC001506	6.8	3.72	2.93	7.69	8.55	5.52	2.68	3.49	3.49								
OVARC001509	1.35	1.98	1.98	5.58	4.61	5.41	3.85	2.97	2.97	**	*	*	*	*	*	*	*
OVARC001510	1.71	1.36	1.70	2.6	2.03	1.38	0.87	1.95	1.95								
OVARC001516	4.33	2.50	2.28	4.35	4.41	5.42	3.48	4.66	4.66								
OVARC001525	1.15	0.47	0.25	2.5	1.57	2.82	0.67	0.87	0.87	*	*	*	*	*	*	*	*
OVARC001543	4.12	4.17	4.21	9.27	7.88	7.75	6.26	7.55	7.55	**	*	*	*	*	*	*	*
OVARC001544	5.06	4.31	3.88	10.18	11.66	10.16	4.91	5.47	5.47	**	*	*	*	*	*	*	*
OVARC001546	4.58	2.37	2.41	4.24	4.04	3.16	5.16	4.17	4.17								
OVARC001547	3.14	1.52	1.67	3.77	5.22	4.03	1.93	2.91	2.91	*	*	*	*	*	*	*	*

128/180

【0769】

【表 298】

1735

1736

QVARC1001916	6.98	5.21	4.19	6.6	6.42	9.56	6.23	7.95	7.95										
QVARC1001928	2.06	0.85	1.79	2.38	2.75	8.84	3.26	4.05	4.05									**	*
QVARC1001937	3.08	3.56	3.08	6.71	6.67	8.66	8.49	10.57	10.57	**									
QVARC1001940	2.73	1.83	2.29	2.29	3.41	3.46	2.76	3.64	3.64	*									
QVARC1001943	7.33	6.50	6.76	5.22	5.72	6.21	4.66	4.79	4.79	*									
QVARC1001943	10.42	8.83	6.98	5.68	5.99	6.08	6.06	4.31	4.31	*									
QVARC1001949	10.36	7.25	8.90	17.76	16.95	14.34	7.64	7.78	7.78	**									
QVARC1001950	6.51	3.98	3.61	6.85	7.18	5.80	4.70	6.17	6.17										
QVARC1001952	8.93	7.35	6.04	9.34	7.56	8.32	8.80	9.41	9.41										
QVARC1001954	2.25	1.93	2.80	2.22	2.67	3.53	3.43	3.10	3.1								*	*	
QVARC1001963	4.35	4.65	3.70	6.06	6.92	7.14	5.20	5.61	5.61	**							*	*	
QVARC1001983	14.69	9.15	11.07	15.77	13.57	18.65	18.62	19.08	19.08								*	*	
QVARC1001987	4.18	3.62	3.23	5.27	5.35	7.29	5.42	5.22	5.22	*							*	*	
QVARC1001989	4.53	2.66	2.25	6.48	8.72	7.41	3.80	4.09	4.09	*							*	*	
QVARC1001991	10.96	5.93	5.69	2.46	8.32	6.27	7.05	6.60	6.6								*	*	
QVARC1002005	5.4	3.75	4.99	8.51	8.21	8.60	5.67	7.46	7.46	**							*	*	
QVARC1002044	5.75	6.74	4.12	8.85	9.04	10.30	6.19	6.78	6.78	*							*	*	
QVARC1002046	11.4	8.29	10.75	14.32	15.39	13.03	16.29	16.11	16.11	*							*	*	
QVARC1002050	7.01	4.34	4.11	5.04	4.91	6.69	6.80	8.61	8.61								*	*	
QVARC1002058	2.46	2.25	3.14	3.04	3.77	4.08	4.59	3.85	3.85								*	*	
QVARC1002066	3.19	1.93	3.61	3.32	2.98	4.14	5.23	6.90	6.9								*	*	
QVARC1002082	4.87	5.01	3.84	11.38	12.17	13.39	6.27	6.19	6.19	**							*	*	
QVARC1002091	9.15	5.09	5.80	7.51	5.64	6.50	4.50	6.13	6.13								*	*	
QVARC1002092	1.08	0.92	1.01	1.95	2.31	1.47	1.26	2.01	2.01	*							*	*	
QVARC1002093	10.46	8.34	8.22	9.65	10.46	8.69	6.29	9.67	9.67								*	*	
QVARC1002094	3.39	2.34	2.33	2.97	3.73	2.67	2.42	4.62	4.62								*	*	
QVARC1002107	4.25	3.34	3.27	6.5	6.62	9.76	3.44	3.77	3.77	*							*	*	
QVARC1002113	10.9	8.09	8.28	16.78	13.09	25.94	13.30	14.51	14.51								*	*	
QVARC1002126	5.65	6.82	6.95	13.64	10.71	12.11	9.13	8.48	8.48	**							*	*	
QVARC1002127	2.58	2.03	3.02	3.02	3.11	2.31	3.36	3.37	3.37								*	*	
QVARC1002138	2.48	2.26	1.89	3.19	3.39	3.93	1.72	2.13	2.13	**							*	*	
QVARC1002143	1.69	1.30	0.60	1.38	1.56	1.86	1.19	0.95	0.95								*	*	
QVARC1002156	1.66	0.93	0.95	1.52	1.87	1.95	2.12	1.74	1.74								*	*	
QVARC1002158	2.7	2.62	1.87	2.12	2.65	2.44	2.26	2.68	2.68								*	*	
QVARC1002165	7.2	5.63	4.73	11.72	8.43	11.59	6.50	7.88	7.88	*							*	*	
QVARC1002176	8	8.96	7.89	12.99	11.14	15.46	14.15	11.02	11.02	*							*	*	
QVARC1002178	1.22	1.02	1.19	6.91	5.74	6.72	4.31	4.39	4.39	**							*	*	
QVARC1002182	2.89	1.94	1.74	3.43	2.78	3.06	2.40	2.34	2.34								*	*	
QVARC1002185	3.07	1.87	2.74	2.77	3.03	2.27	3.08	3.27	3.27								*	*	
PLACE1000004	4.13	1.50	2.40	4.62	3.84	3.14	1.43	2.34	2.34								*	*	
PLACE1000005	1.35	0.94	1.81	2.1	2.21	3.64	1.75	1.86	1.86								*	*	
PLACE1000006	3.24	3.13	3.46	5.32	4.20	5.06	3.54	4.19	4.19	*							*	*	
PLACE1000007	3.52	1.48	1.95	2.76	2.90	3.15	1.95	2.86	2.86								*	*	
PLACE1000014	4.25	3.03	3.71	8.86	8.24	8.01	5.81	6.21	6.21	**							*	*	
PLACE1000031	2.43	0.83	0.85	3.06	2.75	3.91	2.27	1.91	1.91	*							*	*	
PLACE1000033	1.29	0.90	0.41	1.55	1.06	1.17	1.59	1.10	1.1								*	*	
PLACE1000040	4.49	2.71	2.01	6.89	9.12	6.89	4.66	5.42	5.42	*							*	*	
PLACE1000048	1.6	1.02	1.34	5.06	4.76	4.04	3.48	3.87	3.87	**							*	*	
PLACE1000050	5.68	3.49	4.13	5.18	4.97	6.58	3.56	3.95	3.95								*	*	
PLACE1000061	158.3	101.17	90.85	157.97	122.81	120.53	58.82	94.38	94.38								*	*	
PLACE1000066	24.72	10.10	14.31	13.08	14.83	12.97	11.89	17.52	17.52								*	*	
PLACE1000075	3.77	2.50	2.49	11.38	15.88	19.81	6.47	10.82	10.82	**							*	*	
PLACE1000078	3.4	1.72	2.20	4.82	4.89	6.42	2.94	3.88	3.88	*							*	*	
PLACE1000081	10.27	4.42	4.34	7.73	8.15	4.92	5.06	4.85	4.85								*	*	
PLACE1000086	7.07	5.86	4.84	7.21	7.07	4.90	5.17	6.83	6.83								*	*	
PLACE1000094	3.81	2.40	2.03	2.26	2.48	2.45	2.38	2.04	2.04								*	*	

1737

1738

PLACE100101	2.3	2.12	2.61	4.62	5.45	5.31	2.54	3.96	3.96	**	+		
PLACE100121	3.32	1.82	3.36	3.18	3.46	3.22	4.10	2.97	2.97				
PLACE100133	22.32	10.62	12.41	24.57	19.93	22.03	9.44	17.41	17.41				
PLACE100142	3.77	2.94	3.97	3.72	2.78	3.50	4.86	3.02	3.02				
PLACE100146	12.04	5.71	7.52	11.96	8.63	12.12	6.11	6.64	6.64				
PLACE100163	10.38	6.77	6.39	8.08	8.26	4.88	8.20	6.01	6.01				
PLACE100172	2.38	1.36	0.47	1.68	3.26	0.78	1.42	1.36	1.36				
PLACE100181	4.66	3.09	3.18	5.69	5.41	5.62	4.15	4.68	4.68	*	+		
PLACE100184	1.13	1.00	1.41	4.73	6.35	6.17	4.40	7.01	7.01	**	+	**	+
PLACE100185	5.78	3.85	4.83	5.4	6.28	6.49	6.72	6.56	6.56		*	+	+
PLACE100198	3.55	2.09	2.55	2.87	3.21	4.22	3.14	3.19	3.19				
PLACE100213	2.64	0.86	1.73	2.98	2.54	2.75	2.24	2.31	2.31				
PLACE100214	5.38	1.32	2.03	4.15	4.37	5.82	3.05	3.50	3.5				
PLACE100220	5.9	3.44	1.89	3.73	2.84	3.93	2.23	3.16	3.16				
PLACE100231	18.42	11.77	9.30	14.94	14.15	14.87	11.91	14.48	14.48				
PLACE100236	5.6	2.94	3.19	6.04	6.27	4.87	5.66	5.87	5.87				
PLACE100245	7.5	5.11	6.34	10.03	9.79	11.42	4.16	7.99	7.99	**	+		
PLACE100246	5.62	3.38	4.68	6.48	8.30	6.53	8.63	9.43	9.43	*	+	**	+
PLACE100258	15.61	9.21	10.26	23.89	20.68	20.66	9.91	13.07	13.07	*	+		
PLACE100288	2.41	2.18	2.21	2.88	1.68	2.31	2.41	3.07	3.07				
PLACE100292	5.99	4.40	5.17	20.8	17.62	19.45	12.37	20.25	20.25	**	+	**	+
PLACE100302	1.46	1.42	1.22	6.15	8.89	5.78	5.17	5.07	5.07	**	+	**	+
PLACE100304	4.47	1.71	1.91	3.89	2.76	3.12	2.80	2.80	2.8				
PLACE100308	4.91	2.41	1.59	3.39	5.24	3.59	2.01	2.78	2.78				
PLACE100309	11.75	7.68	5.52	7.14	11.13	6.51	7.34	11.09	11.09				
PLACE100313	4.15	1.12	1.95	3.37	3.51	3.75	2.70	2.85	2.85				
PLACE100330	2.07	1.35	1.92	2.05	1.50	2.72	2.22	2.82	2.82		*	+	
PLACE100332	0.54	0.37	0.59	1.08	1.22	2.14	1.43	1.37	1.37	*	+	**	+
PLACE100347	3.56	1.98	2.82	5.26	6.11	4.66	4.59	5.16	5.16	*	+	*	+
PLACE100351	5.67	4.34	5.42	8.3	7.13	5.49	4.92	6.31	6.31				
PLACE100374	9.15	6.32	6.28	12.33	8.13	8.69	5.60	5.63	5.63				
PLACE100380	8.21	2.59	3.63	4.88	6.57	5.07	4.83	5.60	5.6				
PLACE100383	3.43	2.31	1.31	2.37	3.17	2.14	2.59	1.96	1.96				
PLACE100397	4.72	2.15	2.60	3.29	2.51	3.41	2.52	3.33	3.33				
PLACE100401	8.18	4.62	4.15	5.55	6.29	6.94	5.61	6.88	6.88				
PLACE100406	5.56	3.08	2.60	5.54	5.34	5.46	3.82	3.45	3.45				
PLACE100412	3.31	2.01	1.64	4.18	4.67	3.93	2.55	2.54	2.54	*	+		
PLACE100420	10.38	5.91	5.93	8.64	10.82	10.12	5.86	5.89	5.89				
PLACE100421	3.59	3.04	2.31	4.45	4.08	3.36	2.89	3.75	3.75				
PLACE100423	2.95	2.15	1.93	20.49	20.83	20.84	13.81	14.04	14.04	**	+	**	+
PLACE100424	3	2.12	1.66	4.43	3.32	3.59	1.60	2.52	2.52	*	+		
PLACE100430	3.63	1.51	1.58	2.45	2.43	3.11	1.57	3.03	3.03				
PLACE100433	4.59	1.89	2.39	2.55	2.63	3.39	3.84	2.91	2.91				
PLACE100435	4.53	3.13	3.03	9.09	8.39	8.75	5.45	3.19	3.19	**	+		
PLACE100437	2.55	2.34	2.51	7.65	7.02	9.50	7.52	8.64	8.64	**	+	**	+
PLACE100442	12.33	5.94	10.64	23.09	26.07	18.20	10.78	10.42	10.42	*	+		
PLACE100444	9.31	6.03	5.50	16.99	19.05	17.38	8.02	10.01	10.01	**	+		
PLACE100453	6.66	4.79	5.00	7.58	6.74	7.37	6.15	9.05	9.05				
PLACE100456	4.25	3.10	2.24	3.67	3.13	4.02	3.33	4.78	4.78				
PLACE100465	5.73	3.62	3.38	4.99	3.38	5.47	5.67	4.76	4.76				
PLACE100481	5.42	4.78	5.17	5.8	8.48	10.90	5.11	5.58	5.58				
PLACE100492	4.42	2.55	3.57	3.46	5.78	6.28	4.30	4.90	4.9				
PLACE100508	4.11	3.53	2.58	3.28	3.70	3.99	2.37	3.96	3.96				
PLACE100512	5.22	2.40	1.36	6.14	5.78	4.97	4.87	4.74	4.74				
PLACE100540	2.6	2.41	1.99	4.78	4.15	4.34	1.97	1.70	1.7	**	+		
PLACE100541	6.4	6.38	5.54	8.78	7.96	6.93	7.44	11.12	11.12	*	+	*	+

131/180

[0772]

[表301]

1739

1740

PLACE1000546	3.29	1.94	2.26	2.05	2.11	2.19	2.74	2.01	2.01										
PLACE1000547	5.79	5.41	5.37	8.99	6.38	9.62	5.74	8.94	8.94	*									
PLACE1000548	3.31	3.53	2.48	3.26	3.84	4.27	3.23	2.77	2.77										
PLACE1000549	5.48	3.54	4.16	6.47	7.13	6.86	5.29	6.77	6.77	*									
PLACE1000550	2.28	2.89	3.32	2.89	4.25	5.04	4.28	3.71	3.71		*								
PLACE1000551	10.76	7.63	6.51	18.65	16.27	17.87	10.12	7.24	7.24	**									
PLACE1000552	7.2	4.11	4.88	9.4	11.04	9.29	6.85	6.39	6.39	*									
PLACE1000553	7.89	4.98	4.13	9.54	8.74	6.18	7.91	6.38	6.38										
PLACE1000554	7.64	7.46	10.08	8.78	8.56	8.98	4.99	7.82	7.82										
PLACE1000555	5.52	4.56	3.15	8.04	7.54	8.14	4.12	5.23	5.23	**									
PLACE1000556	4.13	3.66	3.53	4.62	5.26	5.10	5.59	5.89	5.89	**									
PLACE1000557	3.95	3.19	2.63	4.04	4.12	4.83	3.09	3.87	3.87										
PLACE1000558	1.33	4.36	3.21	2.64	5.18	3.62	3.25	4.05	4.05										
PLACE1000559	3.93	3.49	2.73	5.31	3.91	4.11	4.05	3.66	3.66										
PLACE1000560	2.72	3.21	2.28	6.49	6.56	3.99	3.45	2.66	2.66	*									
PLACE1000561	2.12	1.92	1.69	2.35	1.07	2.86	1.27	1.58	1.58										
PLACE1000562	2.8	1.22	1.84	2.02	2.53	1.75	1.81	4.26	4.26										
PLACE1000563	9.31	7.34	8.14	10.31	10.47	9.31	12.81	14.00	14		**								
PLACE1000564	1.27	0.67	0.99	1.89	1.74	1.74	1.26	1.65	1.65	*									
PLACE1000565	11.24	11.57	11.40	19.1	16.63	21.24	10.14	12.25	12.25	**									
PLACE1000566	1.84	3.33	4.09	6.55	4.54	5.89	4.64	6.19	6.19	*		*							
PLACE1000567	2.94	0.83	1.14	1.67	1.91	1.48	1.97	1.39	1.39										
PLACE1000568	3.04	1.05	2.32	2.9	2.88	3.09	2.84	2.88	2.88										
PLACE1000569	6.27	3.34	3.42	5.4	6.40	3.86	2.84	3.25	3.25										
PLACE1000570	12.36	6.45	8.51	10.43	9.17	13.07	10.01	13.44	13.44										
PLACE1000571	2.38	1.17	1.02	4.52	3.07	2.68	4.21	4.95	4.95			**							
PLACE1000572	2.51	1.55	1.57	3.46	3.45	4.83	2.60	2.77	2.77	*									
PLACE1000573	2.21	1.01	1.04	2.25	2.24	3.89	2.18	2.07	2.07										
PLACE1000574	5.1	3.19	2.79	4.88	3.83	3.91	3.55	2.38	2.38										
PLACE1000575	8.86	6.54	5.09	10.87	11.53	8.38	4.96	7.33	7.33										
PLACE1000576	4.27	4.46	2.71	4.67	3.49	4.67	3.76	4.74	4.74										
PLACE1000577	6.19	3.54	4.79	9.71	9.92	9.47	5.31	5.48	5.48	**									
PLACE1000578	9.72	4.72	5.55	4.52	4.48	3.39	4.32	4.67	4.67										
PLACE1000579	1.9	1.59	2.33	3.4	3.26	3.47	1.64	2.26	2.26	**									
PLACE1000580	2.3	2.38	1.85	3.32	3.27	4.96	2.41	3.24	3.24	*									
PLACE1000581	7.01	4.40	5.61	12.77	10.75	11.18	7.00	5.92	5.92	**									
PLACE1000582	6.13	3.73	3.27	7.05	6.77	5.20	4.28	5.79	5.79										
PLACE1000583	5.14	3.45	2.78	6.34	7.02	4.42	12.05	18.19	18.19		**								
PLACE1000584	3.14	5.34	2.01	3.49	3.92	2.49	3.35	1.76	1.76										
PLACE1000585	4.46	2.15	3.63	4.5	6.77	4.11	1.87	4.89	4.89										
PLACE1000586	10.82	6.77	8.57	8.51	10.69	9.82	7.58	11.02	11.02										
PLACE1000587	2.83	1.51	2.02	3.37	2.62	2.73	2.59	1.96	1.96										
PLACE1000588	9.64	6.13	6.86	5.2	5.82	6.39	5.18	5.81	5.81										
PLACE1000589	7.89	4.38	5.88	7.14	5.51	8.48	7.94	7.18	7.18										
PLACE1000590	3.08	2.81	1.69	4.08	4.67	3.67	3.31	2.41	2.41	*									
PLACE1000591	16.44	10.14	7.86	22.19	25.12	16.66	7.95	11.86	11.86										
PLACE1000592	3.62	1.21	1.15	2.54	4.35	1.92	1.98	2.37	2.37										
PLACE1000593	6.9	3.41	4.10	5.35	5.89	5.24	4.38	4.49	4.49										
PLACE1000594	3.46	1.48	2.11	2.59	3.24	2.71	3.41	2.78	2.78										
PLACE1000595	0.79	0.41	0.85	0.84	1.40	1.52	0.67	1.33	1.33										
PLACE1000596	3.51	2.64	4.51	6.98	7.67	10.76	8.88	8.80	8.8	*		**							
PLACE1000597	2.76	1.60	7.19	4.08	3.69	6.22	3.38	2.86	2.86										
PLACE1000598	2.02	1.08	0.51	4.48	5.07	3.55	3.07	1.96	1.96	**									
PLACE1000599	3.27	0.90	1.90	2.66	2.46	1.89	1.91	1.97	1.97										
PLACE1000600	2.75	1.53	1.51	2.98	2.99	3.11	3.17	4.29	4.29		*								
PLACE1000601	6.67	4.02	6.08	7.27	8.73	6.46	4.61	7.73	7.73										

132/180

【0773】

【表302】

1743

1744

FLACE1001344	2.76	1.50	1.35	2.41	3.45	2.46	1.70	2.00	2										
FLACE1001351	3.23	1.94	2.24	3.49	3.29	3.25	2.62	4.03	4.03										
FLACE1001366	4.38	2.83	2.63	5.26	5.03	5.59	4.18	3.48	3.48	*	+								
FLACE1001377	2.21	0.95	1.13	1.75	2.13	2.07	1.20	1.68	1.68										
FLACE1001383	3.71	1.90	1.47	3.95	6.26	1.71	1.64	2.49	2.49										
FLACE1001384	3.18	2.05	1.78	4.94	5.31	4.83	2.21	2.83	2.83	**	+								
FLACE1001387	4.38	2.11	2.54	3.04	2.86	4.24	2.34	3.05	3.05										
FLACE1001395	1.59	1.26	1.15	3.65	3.08	5.18	3.82	2.99	2.99	*	+	**	+						
FLACE1001399	11.87	6.31	8.20	17.43	15.28	22.75	13.01	12.96	12.96	*	+								
FLACE1001401	1.52	0.25	1.01	1.14	0.80	1.79	1.18	1.33	1.33										
FLACE1001407	6.8	4.32	5.87	3.76	3.93	5.36	10.73	10.24	10.24			**	+						
FLACE1001412	5.12	1.76	2.22	3.71	2.25	2.65	2.13	1.31	1.31										
FLACE1001414	15.81	9.44	8.70	18.11	13.15	13.80	12.97	12.27	12.27										
FLACE1001416	4.85	3.13	3.24	4.86	3.47	4.68	3.85	4.04	4.04										
FLACE1001433	34.75	27.32	25.94	41.44	46.72	44.79	20.21	24.82	24.82	**	+								
FLACE1001440	3.36	1.52	3.50	3.58	3.41	4.36	3.30	2.97	2.97										
FLACE1001456	2.82	2.23	1.05	4.35	4.43	4.37	3.77	3.38	3.38	*	+								
FLACE1001464	1.12	0.36	0.61	1.11	1.20	1.53	4.05	3.36	3.36			**	+						
FLACE1001468	1	1.48	0.93	1.65	1.22	1.79	1.02	0.92	0.92										
FLACE1001484	5.54	3.35	3.73	7.43	7.35	10.20	3.71	4.16	4.16	*	+								
FLACE1001500	8.54	6.02	4.38	7.39	7.18	5.61	5.36	6.08	6.08										
FLACE1001502	6.06	4.35	3.12	4.46	5.05	6.69	4.11	4.84	4.84										
FLACE1001503	6.09	4.19	3.41	7.11	7.79	6.61	4.97	5.70	5.70	*	+								
FLACE1001505	20.88	12.93	14.68	15.96	17.98	17.32	9.52	14.48	14.48										
FLACE1001513	6.48	3.77	5.22	5.72	3.68	4.54	4.27	6.65	6.65										
FLACE1001516	10.93	7.17	9.57	12.22	8.39	12.84	8.43	11.33	11.33										
FLACE1001517	5.77	3.37	4.96	3.77	4.67	6.00	5.80	4.89	4.89										
FLACE1001523	23.41	10.77	16.66	12.24	9.55	12.27	10.99	12.94	12.94										
FLACE1001526	7.32	4.41	2.62	6.04	11.01	4.64	4.47	5.72	5.72										
FLACE1001534	4	1.96	2.04	4.38	6.28	3.78	3.64	3.03	3.03										
FLACE1001536	2.83	1.23	1.62	1.78	3.23	2.47	3.23	1.81	1.81										
FLACE1001545	36.23	12.22	23.79	37.99	57.83	39.02	33.62	43.32	43.32										
FLACE1001551	6.66	3.51	3.07	3.77	5.41	4.65	3.22	3.12	3.12										
FLACE1001564	1.35	0.83	1.14	1.76	1.17	1.28	1.94	2.02	2.02			**	+						
FLACE1001570	0.93	0.34	0.64	2.16	2.60	4.80	1.89	2.31	2.31	*	+	**	+						
FLACE1001571	7.95	4.12	4.74	8.82	11.30	11.21	6.14	8.15	8.15	*	+								
FLACE1001595	11.86	8.35	6.84	10.13	8.39	8.08	8.16	6.97	6.97										
FLACE1001603	10.71	5.17	5.52	7.59	10.40	7.10	3.81	6.12	6.12										
FLACE1001601	2.7	2.04	2.99	5.01	5.83	4.53	3.42	3.10	3.10	**	+								
FLACE1001608	2.44	2.10	2.41	3.4	4.03	5.05	2.95	3.88	3.88	*	+	*	+						
FLACE1001610	5.43	4.80	5.73	13.88	9.92	13.14	7.65	8.25	8.25	**	+	**	+						
FLACE1001611	3.56	2.47	3.24	3.84	3.75	5.73	3.92	3.82	3.82										
FLACE1001629	6.48	3.78	4.26	6.9	3.97	6.33	1.49	1.62	1.62										
FLACE1001632	8.49	4.09	6.12	10.06	10.02	12.44	6.25	8.12	8.12	*	+								
FLACE1001634	3.06	1.40	1.54	5.61	6.76	4.47	2.48	4.23	4.23	*	+								
FLACE1001637	4.89	3.35	2.51	2.97	3.72	3.26	3.61	4.38	4.38										
FLACE1001640	6.92	2.46	2.49	7.67	8.87	5.84	4.54	7.69	7.69										
FLACE1001655	3.46	2.83	2.76	2.95	2.85	2.93	2.12	2.26	2.26			*	-						
FLACE1001672	3.35	1.68	2.29	4.35	2.93	3.76	4.49	2.60	2.6										
FLACE1001676	1.74	0.78	2.18	1.12	1.17	2.29	1.50	2.79	2.79										
FLACE1001683	8.62	6.71	9.02	11.78	10.73	12.96	10.63	12.86	12.86	*	+	*	+						
FLACE1001691	5.26	3.63	4.10	12.07	10.05	6.33	3.77	5.16	5.16	*	+								
FLACE1001692	4.42	1.12	2.27	4.86	5.36	4.90	4.07	3.26	3.26										
FLACE1001705	8.07	4.26	3.08	6.53	6.17	7.84	4.62	7.50	7.5										
FLACE1001710	3.38	1.68	2.70	3.78	4.88	3.53	4.66	5.71	5.71			*	+						
FLACE1001726	1.91	1.29	2.34	3.39	3.22	2.45	2.24	3.40	3.4	*	+								

134/180

【0775】

【表 3 0 4】

1745

1746

PLACE1001728	1.3	1.02	0.69	1.1	0.60	1.41	1.40	1.39	1.39										
PLACE1001729	6.79	3.57	3.61	3.84	3.10	4.27	2.54	6.08	6.08										
PLACE1001732	9.94	5.41	6.00	8.04	5.84	6.73	6.37	6.11	6.11										
PLACE1001740	1.57	0.32	0.49	0.97	1.11	1.42	1.06	0.82	0.82										
PLACE1001745	5.8	3.72	3.68	4.06	4.33	4.47	4.22	4.88	4.88										
PLACE1001746	3.57	1.52	1.71	4.99	5.18	6.01	3.66	5.62	5.62	*	+	*	+	*	+	*	+	*	+
PLACE1001748	4.5	2.90	2.37	5.53	4.76	3.57	3.80	5.19	5.19										
PLACE1001753	3.51	2.28	3.04	2.88	3.35	3.77	3.11	5.17	5.17										
PLACE1001756	12.16	6.46	7.86	8.59	7.90	8.09	4.55	8.41	8.41										
PLACE1001760	8.72	4.93	5.18	11.47	11.77	9.41	7.48	10.20	10.2	*	+								
PLACE1001767	6.27	4.18	2.75	5.86	5.81	6.64	5.16	5.97	5.97										
PLACE1001771	1.84	1.98	1.82	2.36	2.85	5.41	2.31	1.87	1.87										
PLACE1001775	1.14	0.68	0.37	2.02	1.85	1.82	2.01	0.97	0.97	**	+								
PLACE1001777	17.14	13.64	18.62	21.05	26.38	21.12	40.01	76.23	76.23			*	+						
PLACE1001781	2.45	1.71	2.59	2.44	2.81	2.52	2.91	5.33	5.33										
PLACE1001783	4.43	2.58	2.66	2.32	3.33	2.65	2.54	4.19	4.19										
PLACE1001786	1.74	1.05	1.30	1.23	1.66	1.40	1.26	1.69	1.69										
PLACE1001788	5.13	2.94	2.51	5.8	4.90	5.17	4.40	3.27	3.27										
PLACE1001795	2.72	1.91	2.58	4.69	4.12	5.43	5.56	6.85	6.85	**	+	**	+	**	+	**	+	**	+
PLACE1001799	3.74	3.45	3.29	3.65	3.39	3.75	3.22	5.05	5.05										
PLACE1001810	2.43	0.99	1.08	2.55	2.52	2.29	2.26	1.22	1.22										
PLACE1001817	6.6	4.05	4.21	9.77	8.48	6.29	8.47	8.36	8.36			*	+						
PLACE1001821	3.26	2.45	2.55	4.22	4.44	5.51	4.69	7.27	7.27	*	+	*	+	*	+	*	+	*	+
PLACE1001836	4.29	2.26	1.81	2.56	3.00	3.57	2.41	2.93	2.93										
PLACE1001844	1.78	2.16	1.61	2.8	3.57	4.27	2.87	4.20	4.2	*	+	*	+	*	+	*	+	*	+
PLACE1001845	2.41	1.41	2.18	4.39	5.00	4.06	2.82	2.33	2.33	**	+	**	+	**	+	**	+	**	+
PLACE1001858	4.51	4.42	4.15	7.53	6.22	8.84	4.27	3.55	3.55	*	+	*	+	*	+	*	+	*	+
PLACE1001869	3.09	2.60	2.08	2.74	2.72	3.73	1.99	3.40	3.4										
PLACE1001890	2.77	2.42	1.39	7.46	6.18	5.66	5.49	5.13	5.13	**	+	**	+	**	+	**	+	**	+
PLACE1001897	2.18	2.26	1.85	6.69	5.35	5.34	8.97	9.82	9.82	**	+	**	+	**	+	**	+	**	+
PLACE1001900	31.17	17.00	21.61	32.58	37.84	31.63	15.20	15.90	15.9										
PLACE1001904	3.92	3.02	3.25	2.81	3.73	3.19	4.96	4.49	4.49			*	+						
PLACE1001907	5.11	3.84	3.69	6.62	6.43	7.96	4.32	5.12	5.12	*	+	*	+	*	+	*	+	*	+
PLACE1001910	1.87	3.06	2.35	3.3	3.81	3.68	14.39	26.30	26.3	*	+	**	+	**	+	**	+	**	+
PLACE1001912	2.63	0.79	1.20	4.38	3.77	3.71	2.02	2.67	2.67	*	+	*	+	*	+	*	+	*	+
PLACE1001918	10.38	7.15	8.90	11.66	9.53	15.16	10.15	14.11	14.11										
PLACE1001920	2.53	1.11	1.05	1.68	3.07	1.48	1.79	0.84	0.84										
PLACE1001928	8.17	4.57	3.74	7.72	5.90	6.65	3.44	4.51	4.51										
PLACE1001930	2.19	1.43	2.13	1.81	3.19	3.67	2.17	2.30	2.3										
PLACE1001949	2.08	1.14	1.41	2.07	1.98	1.77	1.69	2.05	2.05										
PLACE1001959	1.52	1.78	2.06	2.37	1.77	2.84	1.64	2.36	2.36										
PLACE1001969	4.16	2.19	2.62	4.17	4.18	4.94	2.88	2.78	2.78										
PLACE1001974	9.4	3.65	4.39	13.34	9.00	13.23	6.71	10.90	10.9										
PLACE1001991	1.69	1.37	1.20	2.64	1.90	2.12	1.52	1.67	1.67	*	+								
PLACE1001993	5.62	5.76	3.72	4.29	4.58	4.62	6.62	4.70	4.7										
PLACE1001999	5.11	2.90	3.88	7.82	6.79	6.73	3.99	4.04	4.04	*	+								
PLACE1002004	8.3	4.91	5.56	11.8	13.71	13.04	6.33	7.42	7.42	**	+								
PLACE1002008	14.39	6.47	3.72	18.67	18.94	18.14	8.81	8.95	8.95	*	+								
PLACE1002015	8.41	6.18	4.44	7.71	7.18	8.85	8.17	7.96	7.96										
PLACE1002044	1.09	1.29	1.71	3.03	2.76	2.81	3.27	3.06	3.06	**	+	**	+	**	+	**	+	**	+
PLACE1002046	3.04	2.60	2.80	3.24	2.46	4.89	3.21	2.77	2.77										
PLACE1002052	1.9	0.59	1.24	2.33	1.00	2.14	1.49	1.25	1.25										
PLACE1002066	6.22	4.18	2.39	10.6	9.81	10.78	7.57	8.32	8.32	**	+	*	+	*	+	*	+	*	+
PLACE1002072	4.3	3.28	3.34	7.74	8.18	6.54	4.13	5.29	5.29	**	+								
PLACE1002073	4.41	2.63	2.33	3.9	3.18	3.49	2.87	3.67	3.67										
PLACE1002080	9.31	4.83	4.67	8.96	9.64	10.72	7.21	6.98	6.98										

135/180

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【表305】

1747

1748

PLACE1002881	1.99	0.89	1.77	2.72	4.23	2.35	2.10	2.07	2.07				
PLACE1002890	14.44	6.66	9.78	10.42	12.14	11.62	5.32	7.78	7.78				
PLACE1002895	6.66	3.83	6.14	8.67	7.29	9.40	5.73	7.69	7.69				
PLACE1002102	11.71	6.09	6.01	11.63	6.93	8.62	6.39	8.11	8.11				
PLACE1002109	2.46	1.22	1.40	2.6	4.68	2.17	2.82	2.11	2.11				
PLACE1002115	3.01	0.88	0.58	1.13	2.98	1.33	0.18	1.10	1.1				
PLACE1002119	18.69	14.15	17.17	28.94	38.25	31.55	24.25	29.45	29.45	**	+	**	+
PLACE1002140	7.37	4.29	6.46	6.39	6.75	7.33	4.91	5.86	5.86				
PLACE1002150	2.02	1.18	2.19	3.93	4.63	3.78	3.27	2.55	2.55	**	+		
PLACE1002153	6.36	3.80	4.46	7.01	6.47	4.93	5.54	4.93	4.93				
PLACE1002157	2.68	1.47	1.39	4.12	3.06	4.68	2.90	3.69	3.69	*	+	*	+
PLACE1002163	7.63	2.62	3.61	7.02	7.14	5.85	5.07	6.08	6.08				
PLACE1002168	4.33	2.82	2.86	4.8	4.18	3.05	4.14	4.00	4				
PLACE1002170	2.98	1.54	1.88	1.56	1.84	1.46	1.96	1.92	1.92				
PLACE1002171	13.45	7.42	8.57	6.89	9.10	5.13	2.02	3.14	3.14		*	-	
PLACE1002180	1.81	0.89	1.51	3.13	3.65	3.26	1.39	2.44	2.44	**	+		
PLACE1002184	2.38	1.68	1.24	6.52	7.00	7.36	6.04	5.01	5.01	**	+	**	+
PLACE1002280	3.74	3.15	2.61	3.65	2.78	3.93	3.98	4.06	4.06				
PLACE1002285	1.24	0.51	0.69	2.33	2.64	4.75	1.98	1.74	1.74	*	+	*	+
PLACE1002213	8.87	4.30	5.26	10.21	8.63	11.56	6.15	7.84	7.84				
PLACE1002219	1.89	0.82	0.74	1.44	2.66	1.62	0.97	0.77	0.77				
PLACE1002227	4.82	2.81	1.66	4.34	4.54	4.85	2.92	3.36	3.36				
PLACE1002253	3.86	2.60	1.93	1.41	2.78	1.93	2.88	2.14	2.14				
PLACE1002256	1.83	0.92	1.11	2.87	3.97	2.85	1.91	3.59	3.59	*	+	*	+
PLACE1002259	3.19	1.70	1.57	6.62	7.59	6.60	5.13	4.07	4.07	**	+	*	+
PLACE1002285	1.77	0.92	0.70	2.37	1.34	1.10	1.30	2.28	2.28				
PLACE1002301	3.7	3.54	3.53	4.57	5.90	8.65	6.82	8.88	8.88		**	+	
PLACE1002310	2.48	1.29	1.37	3.99	3.09	4.29	7.69	9.72	9.72	*	+	**	+
PLACE1002311	3.44	2.13	1.55	3.07	3.48	2.34	2.76	2.43	2.43				
PLACE1002319	4.6	2.18	2.82	2.38	2.25	2.70	1.39	2.13	2.13				
PLACE1002329	4.19	2.99	2.11	3.47	3.41	5.33	3.47	4.66	4.66				
PLACE1002333	1.41	1.34	1.43	2.55	1.71	1.03	1.08	1.25	1.25		*	-	
PLACE1002342	3.55	2.39	2.93	7.53	5.67	7.31	3.57	4.06	4.06	**	+		
PLACE1002343	3.11	2.65	3.16	2.86	3.12	2.88	2.90	5.44	5.44				
PLACE1002355	3.89	1.69	1.70	3.76	3.03	3.60	3.29	2.58	2.58				
PLACE1002358	3.55	2.39	2.49	3.8	3.99	2.81	2.23	2.70	2.7				
PLACE1002359	8	4.42	4.71	3.91	5.64	5.32	4.07	5.01	5.01				
PLACE1002374	14.74	8.20	8.86	9.64	10.72	8.98	11.09	14.20	14.2				
PLACE1002376	7.57	5.16	5.69	9.15	8.50	11.00	8.02	8.55	8.55	*	+	*	+
PLACE1002379	3.61	3.25	3.56	3.36	3.66	3.11	4.20	4.20	4.2		**	+	
PLACE1002386	5.82	2.32	2.77	4.29	2.48	5.32	6.23	7.32	7.32		*	+	
PLACE1002395	5.61	3.00	2.85	4.9	4.34	4.62	4.54	4.04	4.04				
PLACE1002399	2.61	1.20	1.56	3.06	2.87	4.76	3.56	3.31	3.31		*	+	
PLACE1002407	4.59	2.71	2.96	2.81	2.75	3.28	1.95	2.26	2.26				
PLACE1002433	5.13	3.15	3.02	4.68	5.35	5.03	2.23	2.89	2.89				
PLACE1002437	3.54	1.57	2.70	3.76	3.24	3.17	2.59	4.55	4.55				
PLACE1002438	1.21	1.24	1.22	1.63	1.79	2.23	2.23	2.71	2.71	*	+	**	+
PLACE1002446	5.14	2.19	2.50	4.51	3.25	4.13	6.69	7.98	7.98		*	+	
PLACE1002447	2.92	2.41	2.19	1.36	3.06	1.99	2.41	2.93	2.93				
PLACE1002450	1.44	2.03	1.72	3.08	3.13	3.49	2.56	2.00	2	**	+		
PLACE1002462	2.28	1.70	1.59	1.95	1.67	3.14	1.58	2.39	2.39				
PLACE1002468	3.1	2.98	2.42	2.1	3.45	3.05	2.13	2.02	2.02		*	-	
PLACE1002474	2.91	2.82	2.76	8.43	9.88	7.40	6.02	7.81	7.81	**	+	**	+
PLACE1002477	8.13	3.74	5.00	11.28	9.50	9.10	9.42	12.59	12.59	*	+	*	+
PLACE1002493	1.9	2.19	1.21	1.77	1.63	1.93	2.11	3.26	3.26				
PLACE1002497	2.74	1.52	2.43	1.73	1.95	2.51	2.01	3.14	3.14				

136/180

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【表306】

1749

1750

PLACE1002499	3.87	1.99	3.01	5.9	5.94	5.28	3.14	5.21	5.21	**	+		
PLACE1002500	3.82	3.46	3.57	5.63	5.50	7.08	4.28	4.54	4.54	**	+	**	+
PLACE1002514	2.68	2.18	1.93	2.67	2.24	2.48	3.81	2.98	2.98			*	+
PLACE1002518	3.35	3.89	3.09	9.93	9.60	8.45	4.26	3.29	3.29	**	+		
PLACE1002529	1.4	1.36	1.04	1.77	2.14	1.22	1.26	1.64	1.64				
PLACE1002532	8.72	6.46	7.19	6.81	6.68	6.18	7.67	10.12	10.12				
PLACE1002536	4.9	1.91	3.90	5.09	4.55	3.56	4.82	3.96	3.96				
PLACE1002537	3.14	1.37	1.42	3.63	3.37	4.11	2.67	3.94	3.94	*	+		
PLACE1002539	3.39	2.92	3.22	4.41	4.54	5.47	3.82	4.68	4.68	*	+	*	+
PLACE1002547	5.53	5.37	5.59	8.39	7.22	9.28	8.89	10.18	10.18	**	+	**	+
PLACE1002571	4.43	2.94	4.05	4.84	4.88	7.44	3.32	5.08	5.08				
PLACE1002578	5.19	3.96	3.76	12.25	10.98	12.86	5.35	7.25	7.25	**	+	*	+
PLACE1002583	1.66	0.32	1.44	1.04	1.08	1.16	1.18	0.97	0.97				
PLACE1002591	3.86	2.09	2.10	2.84	2.83	2.65	2.44	2.62	2.62				
PLACE1002598	3.84	2.11	2.49	1.35	1.31	2.14	2.05	2.70	2.7				
PLACE1002604	2.65	2.17	1.64	2.8	3.94	3.24	2.43	2.54	2.54				
PLACE1002612	8.01	6.63	6.63	12.1	11.80	12.23	8.71	11.33	11.33	**	+	*	+
PLACE1002629	2.58	1.69	1.51	2.59	2.61	4.00	1.54	3.25	3.25				
PLACE1002638	2.18	2.76	3.22	4.42	3.44	3.29	2.42	3.06	3.06				
PLACE1002655	3.23	4.16	4.18	10.46	6.84	7.33	3.29	5.31	5.31	*	+		
PLACE1002665	4.13	3.33	2.98	6.38	8.83	5.64	3.33	3.33	3.33	*	+		
PLACE1002685	5.53	3.42	2.72	4.03	3.03	3.10	2.10	4.59	4.59				
PLACE1002692	8.51	6.44	4.56	11.47	12.80	10.48	5.21	5.98	5.98	*	+		
PLACE1002714	6.78	4.06	3.36	6.88	8.05	6.09	4.05	4.56	4.56				
PLACE1002721	6.84	4.40	5.49	7.7	6.58	8.37	3.72	4.94	4.94				
PLACE1002723	0.74	0.78	0.84	1.77	1.11	1.67	0.84	1.69	1.69	*	+		
PLACE1002724	3.49	5.71	5.81	8.46	6.47	7.89	5.02	5.08	5.08				
PLACE1002754	3.26	2.58	3.14	6.13	6.45	6.35	3.62	4.77	4.77	**	+	*	+
PLACE1002768	3.97	1.25	1.67	3.3	2.50	2.51	2.78	3.00	3				
PLACE1002772	1.33	0.09	0.96	0.92	1.25	1.40	1.29	1.37	1.37				
PLACE1002775	14.42	7.79	9.64	10.55	11.21	12.27	12.36	11.82	11.82				
PLACE1002789	1.98	1.39	1.23	1.94	2.57	3.18	3.07	5.79	5.79			*	+
PLACE1002782	3.02	0.85	1.61	1.99	1.37	3.05	1.65	1.52	1.52				
PLACE1002794	2.49	1.48	2.20	1.75	2.76	3.63	2.18	2.11	2.11				
PLACE1002795	1.27	0.70	0.60	1.08	1.69	1.49	0.76	0.93	0.93				
PLACE1002811	3.67	1.25	0.81	2.9	2.86	1.50	2.33	2.50	2.5				
PLACE1002815	5.44	2.94	2.29	7.27	15.00	10.36	12.84	16.74	16.74	*	+	**	+
PLACE1002816	8.2	3.96	3.92	6.25	6.25	6.46	5.01	6.23	6.23				
PLACE1002822	3.34	1.86	2.08	2.36	4.04	3.46	2.71	3.36	3.36				
PLACE1002833	7.79	2.79	4.10	8.87	10.15	5.51	5.02	6.82	6.82				
PLACE1002834	10.13	4.35	5.31	13.91	16.48	13.43	4.69	8.55	8.55	*	+		
PLACE1002835	10.05	3.42	5.57	6.59	4.99	6.00	6.06	6.48	6.48				
PLACE1002839	1.69	0.45	1.30	1.57	1.51	2.09	0.58	1.24	1.24				
PLACE1002851	0.76	0.42	0.82	8.19	10.42	3.82	2.14	3.56	3.56	*	+	**	+
PLACE1002853	2.74	1.59	0.90	3.41	5.47	2.91	2.59	4.00	4				
PLACE1002881	6.4	4.73	3.84	9.33	10.46	8.15	5.45	5.65	5.65	*	+		
PLACE1002901	24.97	13.25	21.58	24.2	26.27	27.37	19.88	33.42	33.42				
PLACE1002904	1.92	2.64	1.54	1.71	2.53	1.56	1.90	1.68	1.68				
PLACE1002905	3.55	1.81	2.93	4.92	3.38	5.48	3.25	4.03	4.03				
PLACE1002908	3.03	1.68	2.53	3.06	3.13	3.59	2.41	3.70	3.7				
PLACE1002911	17.43	10.47	13.64	12.95	9.50	14.36	14.97	14.68	14.68				
PLACE1002941	4.78	1.82	1.86	3.71	4.48	2.93	2.39	2.15	2.15				
PLACE1002950	9.17	5.50	4.49	8.77	14.38	6.75	5.85	4.90	4.9				
PLACE1002955	30.87	12.83	9.17	24.34	31.40	20.44	28.45	32.62	32.62				
PLACE1002958	12.6	6.07	8.49	13.92	20.27	16.63	20.63	26.24	26.24	*	+	**	+
PLACE1002963	1.57	0.87	0.79	1.15	2.51	1.72	1.24	1.43	1.43				

137/180

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【表307】

(877)

特開 2002-191363

1751

1752

PLACE1002967	5.1	2.51	3.09	6.76	6.45	5.80	4.18	4.06	4.06	*	+		
PLACE1002968	1.23	0.90	0.78	1.96	2.73	1.63	1.67	2.25	2.25	*	+	**	+
PLACE1002976	14.62	6.59	8.58	14.43	17.61	21.24	10.88	15.26	15.26				
PLACE1002991	9.09	3.33	5.17	10.69	10.94	9.19	4.59	4.38	4.38				
PLACE1002993	4.97	3.72	3.40	7.49	6.57	6.94	4.40	4.67	4.67	**	+		
PLACE1002996	4.17	2.53	2.14	3.73	3.53	2.43	2.53	3.20	3.2				
PLACE1003010	14.09	9.21	8.66	11.39	9.01	12.05	11.70	11.85	11.85				
PLACE1003025	3.37	1.92	1.25	3.12	3.46	2.82	2.56	2.83	2.83				
PLACE1003027	2.78	1.30	1.63	3.36	4.14	4.94	2.51	3.33	3.33	*	+		
PLACE1003044	5.29	2.38	3.63	5.05	4.60	4.39	4.30	3.74	3.74				
PLACE1003045	1.31	0.14	0.41	1.12	0.74	1.58	0.92	1.66	1.66				
PLACE1003052	5.81	2.44	2.52	4.24	6.72	5.03	2.74	4.06	4.06				
PLACE1003083	1.98	0.63	0.30	1.59	1.48	1.45	1.09	1.36	1.36				
PLACE1003085	8.86	4.56	4.41	4.48	5.13	3.76	5.79	5.25	5.25				
PLACE1003092	4.95	2.80	2.49	4.61	7.21	5.11	3.15	5.59	5.59				
PLACE1003097	2.48	1.08	1.75	2.13	2.19	3.46	1.83	1.87	1.87				
PLACE1003100	5.55	3.04	3.54	4.48	2.63	4.78	3.66	4.38	4.38				
PLACE1003108	2.43	2.01	1.88	3.79	4.20	5.56	3.02	3.15	3.15	*	+	**	+
PLACE1003115	5.59	4.45	4.08	5.2	3.47	4.38	3.94	4.36	4.36				
PLACE1003120	9.1	5.05	6.99	11.92	11.69	8.39	4.33	5.35	5.35				
PLACE1003135	7.15	3.42	2.81	2	1.71	2.50	1.33	2.53	2.53				
PLACE1003136	9.4	3.19	5.96	7.56	7.72	8.01	6.80	8.18	8.18				
PLACE1003141	1.43	1.20	0.97	1.12	1.71	2.12	1.29	2.62	2.62				
PLACE1003145	1.17	1.88	1.88	1.29	0.85	1.19	1.52	2.74	2.74				
PLACE1003147	3.88	1.84	2.10	3.04	3.09	5.16	2.94	6.44	6.44				
PLACE1003153	2.04	1.22	1.34	1.76	3.27	2.50	1.12	2.13	2.13				
PLACE1003163	5.21	2.54	2.21	3.71	2.70	3.59	1.58	3.29	3.29				
PLACE1003172	17.21	13.29	11.63	20.51	17.81	16.21	12.82	14.76	14.76				
PLACE1003174	1.86	0.95	0.96	2.33	2.68	2.13	2.07	2.85	2.85	*	+	*	+
PLACE1003176	1.87	0.85	0.99	0.69	1.79	1.46	1.77	2.02	2.02				
PLACE1003181	2.42	1.29	1.30	1.36	1.88	1.93	2.33	2.76	2.76				
PLACE1003184	4.02	2.35	1.57	1.09	1.42	1.68	2.02	2.95	2.95				
PLACE1003190	12.59	7.17	8.42	3.7	4.03	4.95	5.55	3.22	3.22	*	-	*	-
PLACE1003200	0.16	0.08	0.11	0.98	0.55	0.76	0.91	1.63	1.63	**	+	**	+
PLACE1003205	10.63	4.75	4.99	13.42	19.02	15.56	5.60	9.62	9.62	*	+		
PLACE1003209	1.33	0.58	0.91	1.06	1.71	1.13	1.44	1.84	1.84		*	+	
PLACE1003214	3.74	1.92	0.96	2.48	3.08	2.07	2.80	1.58	1.58				
PLACE1003229	4.01	2.47	1.89	4.67	6.17	5.71	3.46	3.20	3.2	*	+		
PLACE1003238	0.55	1.29	0.72	1.01	1.04	1.42	1.89	4.82	4.82		*	+	
PLACE1003249	4.21	2.68	2.29	5.89	6.34	7.49	3.21	4.18	4.18	**	+		
PLACE1003256	15.42	10.76	11.86	18.06	20.59	21.48	20.54	17.58	17.58	*	+	*	+
PLACE1003268	1.59	3.70	0.75	1.91	1.78	1.15	1.24	1.39	1.39				
PLACE1003279	5.6	4.25	1.88	7.33	8.87	7.26	3.36	5.26	5.26	*	+		
PLACE1003294	5.96	3.04	2.55	5.19	4.93	5.17	2.65	4.69	4.69				
PLACE1003296	3.69	1.73	1.93	4.06	3.41	2.82	2.94	3.29	3.29				
PLACE1003297	6.38	2.82	3.60	6.92	8.35	6.63	3.36	5.38	5.38				
PLACE1003302	6.92	3.76	5.11	9.34	12.52	9.10	7.08	7.90	7.9	*	+		
PLACE1003334	0.67	1.10	1.68	2.93	3.55	4.00	2.24	3.22	3.22	**	+	*	+
PLACE1003337	10.11	6.39	4.50	7.53	11.58	10.30	6.19	6.17	6.17				
PLACE1003342	1.8	1.54	1.48	1.86	2.85	3.11	2.71	3.37	3.37		**	+	
PLACE1003343	0.54	0.36	0.34	0.71	0.79	1.22	0.47	0.55	0.55	*	+		
PLACE1003344	24.27	18.53	13.01	17.74	21.43	22.39	17.02	18.57	18.57				
PLACE1003353	17.73	10.09	9.18	16.17	17.86	14.13	8.53	9.79	9.79				
PLACE1003361	5.88	2.41	3.54	11.34	11.94	9.89	3.99	5.89	5.89	**	+		
PLACE1003366	6.48	3.29	5.05	6.5	8.30	6.96	4.61	4.25	4.25				
PLACE1003369	2.89	2.16	1.46	3.79	2.79	2.98	2.58	2.98	2.98				

138/180

【0779】

【表308】

(878)

特開2002-191363

1753

1754

PLACE1003373	4.86	3.69	3.10	6.36	6.08	6.40	5.24	6.26	6.26	*	+	*	+
PLACE1003373	4.59	2.14	1.77	6.44	8.87	7.14	3.34	3.58	3.58	*	+		
PLACE1003375	1.64	2.20	2.31	1.72	2.46	2.62	1.19	1.43	1.43			*	-
PLACE1003378	2.12	1.60	1.84	2.23	1.84	1.68	2.18	2.69	2.69				
PLACE1003380	2.45	1.53	0.51	2.22	2.04	0.76	1.14	1.36	1.36				
PLACE1003394	8.16	3.88	4.89	10.77	12.17	8.54	8.17	10.02	10.02	*	+		
PLACE1003401	3.67	0.79	0.99	1.2	1.46	1.82	0.45	1.86	1.86				
PLACE1003405	6.01	6.00	6.98	4.76	7.61	8.04	6.47	7.65	7.65				
PLACE1003407	4.49	4.04	3.71	5.05	5.22	5.15	5.12	5.61	5.61	*	+	**	+
PLACE1003420	4.75	4.07	3.59	7.55	10.89	8.12	4.15	6.01	6.01	*	+		
PLACE1003428	2.19	2.41	3.05	3.29	4.02	4.47	2.16	2.43	2.43	*	+		
PLACE1003432	7.17	3.85	3.68	4.37	7.22	7.66	3.81	6.34	6.34				
PLACE1003438	9.06	3.37	4.39	5.86	7.12	5.43	5.87	7.15	7.15				
PLACE1003452	3.13	1.08	2.21	1.29	5.01	2.29	2.22	2.52	2.52				
PLACE1003454	8.4	4.68	5.18	7.33	6.34	9.17	4.92	7.46	7.46				
PLACE1003455	13.75	5.01	6.05	6.83	8.91	9.83	8.45	9.21	9.21				
PLACE1003456	7.28	4.38	4.13	10.64	12.00	13.60	7.62	7.20	7.2	**	+		
PLACE1003460	7.84	3.76	6.10	10.15	7.44	7.77	6.55	7.66	7.66				
PLACE1003478	3.33	0.56	0.93	2.01	1.78	1.24	0.65	0.96	0.96				
PLACE1003484	7.55	4.57	2.88	11.32	16.35	7.83	7.21	9.47	9.47				
PLACE1003493	14.03	6.96	6.73	11.22	11.97	14.63	9.74	9.34	9.34				
PLACE1003503	42.11	19.93	34.28	29.63	36.26	35.89	25.50	29.49	29.49				
PLACE1003505	2.24	1.06	0.89	0.91	0.90	1.59	2.08	1.73	1.73				
PLACE1003516	1.01	0.49	0.89	2.17	2.40	2.58	1.68	1.86	1.86	**	+	**	+
PLACE1003519	39.78	23.99	30.04	55.6	50.01	57.71	22.97	28.09	28.09	*	+		
PLACE1003520	45.85	22.30	34.27	66.52	30.94	72.87	38.79	44.73	44.73				
PLACE1003521	1.43	0.65	0.89	2.33	3.32	0.95	2.10	3.87	3.87		*	+	
PLACE1003525	15.69	8.19	8.09	12.57	19.45	12.58	15.38	18.26	18.26				
PLACE1003528	126.72	75.71	77.51	102.34	128.72	89.84	56.09	57.39	57.39				
PLACE1003529	10.31	6.25	7.90	10.63	11.63	11.54	9.31	9.78	9.78				
PLACE1003537	3.45	1.76	2.18	3.36	4.60	3.48	5.58	5.15	5.15			**	+
PLACE1003549	3.96	2.80	3.67	4.57	2.88	5.08	2.97	4.32	4.32				
PLACE1003553	6.15	2.35	3.07	4.85	4.12	5.00	3.14	3.29	3.29				
PLACE1003566	5.25	2.36	2.80	5.45	5.03	6.90	4.92	5.27	5.27				
PLACE1003568	1.39	1.43	0.56	1.66	1.56	1.27	1.01	0.83	0.83				
PLACE1003573	2.04	1.89	1.09	2.09	2.81	1.71	1.61	1.69	1.69				
PLACE1003575	3.94	2.36	1.55	4.2	5.03	5.48	3.67	2.41	2.41	*	+		
PLACE1003583	1.25	0.21	0.91	0.63	1.54	1.28	1.19	0.85	0.85				
PLACE1003584	3.17	2.52	1.33	5.76	4.75	5.94	2.30	3.30	3.3	**	+		
PLACE1003592	6.37	4.34	3.44	8.54	12.20	11.57	7.98	8.85	8.85	*	+	*	+
PLACE1003593	0.73	1.09	0.64	1.3	1.69	1.81	0.49	1.57	1.57	*	+		
PLACE1003594	16.13	4.42	11.69	14.87	17.87	21.56	10.51	11.29	11.29				
PLACE1003596	5.64	5.18	5.93	10.49	15.28	7.57	7.20	9.60	9.6		*	+	
PLACE1003598	13.48	8.08	6.25	8.41	8.96	8.69	7.81	8.78	8.78				
PLACE1003602	3.72	2.13	1.45	3.5	3.57	3.37	2.64	3.45	3.45				
PLACE1003605	18.39	10.93	10.02	16.96	17.66	21.30	9.74	14.50	14.5				
PLACE1003611	3.07	0.86	1.19	2.62	2.97	3.49	1.69	2.05	2.05				
PLACE1003618	2.42	0.71	0.96	1.64	1.41	1.56	1.78	2.12	2.12				
PLACE1003625	3.62	1.30	2.39	3.11	4.04	4.15	3.30	3.49	3.49				
PLACE1003626	13.07	5.94	8.16	14.48	13.10	14.74	12.62	11.51	11.51				
PLACE1003630	3.48	2.42	1.94	3.18	3.39	2.97	3.11	3.27	3.27				
PLACE1003635	2.04	1.03	1.44	2.07	2.17	2.34	1.81	1.67	1.67				
PLACE1003638	3.27	2.36	1.79	4.52	4.52	3.82	3.33	3.31	3.31	*	+		
PLACE1003644	3.31	2.33	2.10	5.21	5.95	5.73	4.05	4.05	4.05	**	+	*	+
PLACE1003654	4.23	1.54	1.89	1.81	2.78	2.00	0.89	2.32	2.32				
PLACE1003656	2.23	0.80	1.38	1.4	1.47	1.90	1.48	2.10	2.1				

138/180

【0780】

【表309】

1758

FLACE1004104	2.15	1.27	0.85	1.43	1.39	2.13	1.09	2.01	2.01				
FLACE1004113	4.08	1.68	3.31	4.66	4.46	4.54	3.36	3.05	3.05				
FLACE1004114	2.54	0.84	0.51	1.58	2.53	1.82	2.42	1.88	1.88				
FLACE1004118	1.28	1.29	1.42	1.63	4.01	2.38	1.61	2.11	2.11				
FLACE1004128	12.83	9.07	9.04	8.02	8.50	9.63	5.06	6.17	6.17				
FLACE1004139	2.24	2.05	1.92	1.83	3.44	3.33	1.32	1.72	1.72				
FLACE1004149	18	9.56	12.62	22.09	23.13	25.79	15.85	17.31	17.31	*			
FLACE1004156	8.66	4.78	4.97	11.23	13.14	12.83	5.87	8.14	8.14	*	+		
FLACE1004160	31.97	23.56	27.55	20.37	16.69	25.95	28.83	35.50	35.5				
FLACE1004161	12.19	6.98	6.65	7.81	8.30	9.68	8.49	8.65	8.65				
FLACE1004166	10.59	4.49	3.61	8.56	19.40	8.04	5.20	7.58	7.58				
FLACE1004168	9.22	3.40	4.94	7.74	9.05	6.39	5.52	5.88	5.88				
FLACE1004170	0.56	0.65	1.17	2.02	1.70	2.28	1.72	2.24	2.24	**	+	**	+
FLACE1004178	5.68	2.50	3.59	4.97	6.58	6.01	4.61	7.20	7.2				
FLACE1004183	4.44	2.26	4.45	5.52	5.64	5.63	4.08	3.85	3.85				
FLACE1004197	1.06	1.17	1.74	1.07	1.49	1.13	2.10	1.67	1.67				
FLACE1004199	9.96	4.47	8.63	4.5	6.39	5.99	10.80	9.20	9.2				
FLACE1004203	6.09	3.61	5.37	4.74	4.70	4.68	5.77	5.62	5.62				
FLACE1004241	7.53	2.60	2.25	8.1	9.90	6.46	4.60	5.49	5.49				
FLACE1004249	25.51	14.54	13.20	20.9	26.96	19.21	17.71	21.13	21.13				
FLACE1004255	1.02	0.73	0.36	0.86	1.57	1.36	0.69	1.07	1.07				
FLACE1004256	4.42	1.01	3.09	9.24	13.36	13.94	12.44	10.96	10.96	**	+	**	+
FLACE1004257	4.54	1.21	1.79	4.96	4.55	4.58	3.59	4.84	4.84				
FLACE1004258	3.59	2.38	2.35	2.98	2.70	2.85	3.20	2.02	2.02				
FLACE1004270	3.93	3.24	3.36	3.85	4.28	6.05	3.70	3.05	3.05				
FLACE1004273	4.04	2.85	3.28	3.85	5.74	5.17	3.42	6.23	6.23				
FLACE1004275	83.7	57.27	49.34	101.5	84.19	78.07	49.24	46.63	46.63				
FLACE1004274	2.95	0.92	1.52	1.53	2.26	1.62	1.54	1.70	1.7				
FLACE1004277	4.89	3.63	3.77	5.98	6.33	5.84	3.49	5.35	5.35	*	+		
FLACE1004279	4.14	3.27	2.56	4.12	4.89	5.01	2.41	5.41	5.41				
FLACE1004282	4.87	1.71	2.16	3.7	2.78	3.26	3.33	4.30	4.3				
FLACE1004284	5.6	3.43	5.55	7.94	7.12	9.08	5.18	6.08	6.08	*	+		
FLACE1004289	4.45	2.76	2.32	4.87	4.64	6.03	3.57	3.74	3.74				
FLACE1004299	3.82	1.87	1.73	3.07	2.88	4.42	3.05	2.95	2.95				
FLACE1004302	2.2	0.86	0.90	1.74	3.32	2.03	1.19	1.35	1.35				
FLACE1004305	3.85	2.26	1.99	1.85	1.24	2.43	2.28	2.58	2.58				
FLACE1004316	5.43	2.71	3.07	1.96	3.30	2.21	2.72	4.32	4.32				
FLACE1004323	1.43	0.69	0.73	1.49	2.28	1.46	1.11	2.06	2.06				
FLACE1004325	13.88	6.16	7.35	9.82	9.01	12.35	11.00	10.37	10.37				
FLACE1004332	3.01	1.40	1.75	1.66	1.82	2.98	2.54	3.00	3				
FLACE1004336	5.91	5.69	5.62	10.43	10.12	10.42	6.74	8.77	8.77				
FLACE1004346	3.07	2.03	1.73	2.75	2.78	2.82	1.63	2.50	2.5				
FLACE1004358	17.58	10.51	10.45	12.26	12.55	14.79	12.38	16.11	16.11				
FLACE1004376	21.68	10.31	10.00	13.3	12.08	12.35	12.00	16.69	16.69				
FLACE1004384	3.8	1.61	2.13	4.74	4.70	5.37	3.12	3.81	3.81	*	+		
FLACE1004385	1.9	0.86	0.50	0.57	0.48	1.53	0.60	1.25	1.25				
FLACE1004388	3.6	1.83	1.85	3.69	4.57	4.12	1.57	1.95	1.95				
FLACE1004485	0.61	1.07	0.82	0.21	0.91	1.17	2.14	2.86	2.86			**	+
FLACE1004487	5.17	3.33	4.50	6.41	4.91	7.01	4.80	3.58	3.58				
FLACE1004424	1.66	0.59	0.46	0.66	0.44	2.14	0.23	0.46	0.46				
FLACE1004425	1.47	0.52	1.48	2.94	3.61	2.12	1.72	2.14	2.14	*	+		
FLACE1004427	2.86	1.31	1.07	1.87	2.05	1.81	1.95	3.44	3.44				
FLACE1004428	3.96	2.20	1.76	4.03	5.57	4.86	3.58	3.24	3.24				
FLACE1004433	6.32	3.82	4.97	5.63	4.68	5.19	2.95	4.99	4.99				
FLACE1004435	7.56	3.49	4.09	10.74	10.16	12.36	5.74	11.13	11.13	*	+		
FLACE100437	7.97	3.59	4.68	4.42	7.20	5.02	5.17	3.07	3.07				

【表 3 1 1】

1759

1760

PLACE1004441	3.25	1.90	2.33	4.32	4.15	5.16	3.84	4.52	4.52	*	+	+
PLACE1004446	1.76	2.09	0.72	1.34	1.42	1.87	2.28	2.32	2.32			
PLACE1004450	0.76	0.23	0.38	0.96	1.30	0.99	0.73	0.72	0.72	*	+	
PLACE1004451	2.04	1.05	0.94	1.87	2.71	1.33	1.83	2.40	2.4			
PLACE1004456	13.14	7.90	8.58	15.19	13.06	9.85	9.75	13.11	13.11			
PLACE1004458	1.13	0.48	0.38	2.8	2.09	3.55	9.05	9.62	9.62	*	**	+
PLACE1004460	1.24	0.45	0.57	1.15	1.35	1.69	1.34	1.71	1.71		*	+
PLACE1004467	6.23	3.77	6.46	8.7	9.58	9.65	5.25	4.76	4.76	*	+	
PLACE1004471	7.06	5.28	5.80	10.51	12.81	16.26	6.17	7.08	7.08	*	+	
PLACE1004473	1.57	1.48	1.06	1.91	1.92	2.41	1.84	1.43	1.43	*	+	
PLACE1004475	17.9	8.89	9.13	27.5	24.29	13.71	28.08	20.33	20.33	*	+	
PLACE1004482	2.18	1.39	1.16	1.98	2.90	3.51	2.75	3.78	3.78	*	+	
PLACE1004491	0.74	0.46	0.72	0.47	1.01	0.56	0.69	1.94	1.94			
PLACE1004492	33.34	16.09	17.54	17.67	22.65	21.39	20.85	24.45	24.45			
PLACE1004506	5.1	3.77	3.89	3.53	5.30	4.79	5.63	7.41	7.41	*	+	
PLACE1004507	2.94	1.98	2.25	1.75	2.11	1.80	2.62	3.67	3.67			
PLACE1004510	2.01	2.57	2.33	4.62	4.58	4.58	3.18	2.57	2.57	**	+	
PLACE1004516	1.04	0.43	0.32	0.6	0.82	1.51	0.69	1.14	1.14			
PLACE1004518	5.88	3.35	1.73	3.03	3.63	1.95	4.27	3.46	3.46			
PLACE1004519	3.55	1.36	2.17	1.53	2.33	1.77	1.26	1.42	1.42			
PLACE1004520	4.8	1.73	3.29	3.58	4.49	2.98	3.20	4.60	4.6			
PLACE1004530	7.81	5.59	5.82	2.93	4.17	2.72	3.17	3.36	3.36	*	*	-
PLACE1004545	0.98	1.24	0.71	1.02	1.35	1.28	1.23	1.48	1.48			
PLACE1004547	3.48	2.58	2.62	3.89	3.59	4.14	3.27	6.00	6	*	+	
PLACE1004548	5.32	3.02	2.13	5.34	7.57	7.29	2.74	4.90	4.9			
PLACE1004550	4.75	3.89	2.55	4.32	5.77	4.11	3.73	5.54	5.54			
PLACE1004551	2.21	1.18	1.01	2.32	3.16	1.67	1.47	1.73	1.73			
PLACE1004559	1.69	0.68	1.41	2.2	2.41	1.95	1.58	1.77	1.77	*	+	
PLACE1004562	7.92	4.63	4.61	12.8	13.69	12.24	11.70	16.91	16.91	**	+	+
PLACE1004564	5.08	3.48	2.94	3.43	4.16	2.75	2.50	3.03	3.03			
PLACE1004604	1.61	1.65	0.87	1.96	1.66	1.23	6.31	2.27	2.27			
PLACE1004611	6.51	4.71	3.22	13.38	14.72	11.15	6.91	6.89	6.89	**	+	
PLACE1004629	3.8	3.23	3.16	7.62	7.80	6.85	5.92	7.19	7.19	**	+	+
PLACE1004630	4.43	7.59	4.92	4.3	3.84	5.63	3.88	4.82	4.82	*	+	
PLACE1004637	9.71	8.66	5.16	8.97	5.26	6.98	6.87	7.85	7.85			
PLACE1004645	34.24	15.91	17.01	26.16	30.73	32.52	15.81	17.34	17.34			
PLACE1004646	3.38	1.74	3.32	3.28	4.81	3.28	2.79	2.82	2.82			
PLACE1004648	14.4	8.71	8.36	10.69	11.92	11.82	11.67	15.16	15.16			
PLACE1004655	41.73	23.86	25.42	40	42.96	45.63	19.14	24.74	24.74			
PLACE1004658	4.07	3.17	2.80	4.22	4.91	5.38	4.38	3.84	3.84	*	+	
PLACE1004664	2.14	1.15	0.86	2.2	2.05	3.93	1.74	1.79	1.79			
PLACE1004672	11.36	7.67	9.44	13.22	15.37	20.21	6.56	12.23	12.23	*	+	
PLACE1004674	6.89	4.27	3.73	8.23	11.59	6.63	7.24	9.33	9.33	*	+	
PLACE1004681	5.36	2.49	2.37	3.93	6.34	2.28	3.03	2.81	2.81			
PLACE1004686	4.25	1.52	2.69	8.28	8.25	8.23	3.83	5.37	5.37	**	+	
PLACE1004690	27.35	18.33	25.68	19.28	26.77	23.31	7.55	15.04	15.04	*	-	
PLACE1004691	4.78	2.55	2.69	4.7	7.55	6.34	2.68	5.61	5.61			
PLACE1004693	3.07	1.09	1.84	2.44	2.98	3.35	2.53	3.19	3.19			
PLACE1004701	23.69	11.94	19.76	25.99	21.50	33.00	23.49	24.31	24.31			
PLACE1004705	5.61	4.43	3.93	4.87	5.07	5.49	3.83	4.06	4.06			
PLACE1004708	9.98	7.05	4.96	7.36	17.22	7.73	12.21	12.77	12.77	*	+	
PLACE1004716	5.47	2.91	3.32	5.79	8.69	5.11	4.23	4.07	4.07			
PLACE1004722	1.53	1.35	1.55	1.7	3.73	3.67	0.90	2.30	2.3			
PLACE1004736	16.73	9.74	14.43	13.11	17.99	18.80	11.66	17.15	17.15			
PLACE1004737	2.18	1.67	1.79	1.43	3.14	3.32	1.37	1.80	1.8			
PLACE1004740	6.4	3.17	4.45	6.16	4.57	7.19	5.04	5.92	5.92			

142/180

【0783】

【表312】

1762

FLACE1004743	2.83	1.69	1.62	2.65	2.30	3.35	1.64	2.59	2.59				
FLACE1004751	3.88	2.76	2.71	4.3	4.81	6.07	2.16	4.03	4.03	+			
FLACE1004757	6.62	2.79	3.38	5.64	5.36	5.13	4.59	3.33	3.33				
FLACE1004761	1.53	0.69	0.99	1.89	2.90	1.43	1.17	2.01	2.01				
FLACE1004773	6.07	1.81	3.15	5.28	4.05	5.04	3.00	3.37	3.37				
FLACE1004775	0.59	0.48	0.41	0.54	0.33	0.45	0.35	1.11	1.11				
FLACE1004777	2.87	1.56	1.63	3.6	3.28	3.27	3.12	2.18	2.18	*	+		
FLACE1004793	1.91	0.67	0.75	1.6	1.01	2.08	1.33	1.74	1.74				
FLACE1004796	11.15	4.76	6.53	15.2	11.67	18.12	12.53	11.15	11.15	+			
FLACE1004804	2.49	2.83	3.47	3.45	3.93	5.58	2.84	4.15	4.15				
FLACE1004813	1.83	1.78	1.19	2.06	4.34	2.04	2.93	2.51	2.51			**	+
FLACE1004814	15.6	8.20	7.30	20.97	26.36	22.14	11.65	11.36	11.36	*	+		
FLACE1004815	2.09	1.04	1.32	4.73	4.30	3.36	2.27	2.36	2.36	**	+		
FLACE1004816	3.22	1.11	2.11	2.58	2.27	3.19	1.56	4.07	4.07				
FLACE1004824	10.16	4.47	7.27	17.15	18.66	21.40	8.53	11.08	11.08	**	+		
FLACE1004827	3.25	1.26	2.36	5.76	5.15	4.86	3.26	3.82	3.82	**	+		
FLACE1004836	2.02	0.78	1.32	3.29	3.51	3.51	1.36	2.69	2.69	**	+		
FLACE1004838	3.17	2.09	1.89	2.78	2.46	3.36	1.52	3.28	3.28				
FLACE1004840	1.23	0.56	0.64	2.27	3.76	2.10	1.40	1.24	1.24	*	+		
FLACE1004843	5.48	1.99	1.07	1.39	1.40	2.34	2.69	3.06	3.06				
FLACE1004850	3.11	1.83	1.19	2.34	1.99	1.83	2.00	3.44	3.44				
FLACE1004868	1.78	1.97	1.38	1.05	1.30	0.94	1.18	1.52	1.52	*	-		
FLACE1004885	4.12	2.86	3.03	6.17	4.95	6.21	2.81	3.69	3.69	*	+		
FLACE1004886	1.77	1.59	1.70	1.43	1.55	1.82	3.23	4.30	4.30			*	+
FLACE1004887	25.24	11.67	14.76	21.81	38.02	28.05	8.65	10.31	10.31				
FLACE1004896	2.33	1.72	1.45	4.61	4.55	3.16	5.89	7.01	7.01	*	+	**	+
FLACE1004900	9.03	4.30	5.53	9.31	10.97	9.80	5.74	6.69	6.69				
FLACE1004902	15.98	5.16	8.41	6.64	13.40	8.82	7.56	8.91	8.91				
FLACE1004904	2.63	1.32	1.15	1.84	2.37	1.90	3.74	3.50	3.5			*	+
FLACE1004911	1.14	3.11	1.00	4.23	0.30	0.65	0.27	1.36	1.36				
FLACE1004913	2.14	1.21	1.21	2.7	1.96	3.02	1.97	4.39	4.39				
FLACE1004918	1.11	0.31	1.10	1.32	1.60	1.48	0.91	1.02	1.02				
FLACE1004930	3.51	2.35	1.88	1.71	2.51	2.60	1.12	1.41	1.41				
FLACE1004934	2.04	1.42	1.26	1.7	2.74	2.49	1.45	1.52	1.52				
FLACE1004937	5.11	2.46	1.95	3.63	3.54	3.36	2.75	2.15	2.15				
FLACE1004949	4.03	1.71	2.54	6.88	7.76	8.45	5.04	2.82	2.82	*	+	*	+
FLACE1004966	3.48	2.29	1.51	2.73	3.17	3.01	2.31	4.32	4.32				
FLACE1004970	0.79	0.82	0.40	0.36	1.00	0.91	0.81	2.69	2.69				
FLACE1004972	1.78	1.50	1.56	2.23	2.38	3.07	1.16	2.50	2.5	*	+		
FLACE1004974	3.63	3.03	1.68	3.41	3.31	2.59	1.64	1.70	1.7				
FLACE1004975	4.46	3.12	2.44	4.13	3.11	5.49	3.51	3.95	3.95				
FLACE1004979	4.8	5.17	3.63	8.89	10.47	10.51	5.50	6.33	6.33	**	+	*	+
FLACE1004982	12.69	7.06	8.29	13.78	13.06	8.17	7.03	8.87	8.87				
FLACE1004985	2.12	1.35	0.79	2.05	1.96	1.11	0.99	3.21	3.21				
FLACE1005003	3.67	1.05	1.88	1.3	2.66	1.79	0.59	2.43	2.43				
FLACE1005004	1.24	1.06	1.30	1.55	1.31	1.17	1.68	1.83	1.83			**	+
FLACE1005005	8.08	4.02	3.41	8.61	8.51	8.54	5.01	5.29	5.29				
FLACE1005011	2.2	1.69	2.79	3	3.06	5.33	3.11	2.57	2.57				
FLACE1005026	2.34	1.90	2.06	1.86	2.01	2.93	1.82	1.53	1.53	*	+	-	
FLACE1005027	4.99	3.43	4.26	7.85	11.24	9.53	3.37	5.57	5.57	*			
FLACE1005031	6.43	2.62	2.97	5.45	4.09	3.20	3.04	3.84	3.84				
FLACE1005036	7.51	3.86	5.10	9.33	12.02	7.99	3.66	4.98	4.98				
FLACE1005041	0.87	0.62	0.58	1.76	1.87	1.43	1.58	1.91	1.91	**	+	**	+
FLACE1005046	7.09	3.32	3.54	11.11	10.13	10.18	4.94	5.99	5.99	**	+		
FLACE1005047	3.57	1.97	1.47	3.2	3.87	3.39	2.49	3.04	3.04				
FLACE1005052	4.36	2.90	3.32	3.11	2.49	4.07	4.21	4.75	4.75				

1763

1764

PLACE1005055	1.93	1.90	2.25	2.55	3.80	3.83	1.39	2.30	2.3	*	+		
PLACE1005066	3.73	3.53	2.95	3.62	2.74	3.71	4.65	6.92	6.92		*	+	
PLACE1005077	1.88	0.74	0.51	1.94	2.30	1.62	1.19	1.27	1.27				
PLACE1005085	5.35	2.26	1.94	7.82	9.01	6.89	4.04	4.10	4.1	*	+		
PLACE1005086	8.18	4.09	4.61	8.82	11.72	8.88	4.94	5.91	5.91				
PLACE1005088	48.83	27.68	29.69	27.61	39.82	34.65	26.01	25.68	25.68				
PLACE1005089	2.42	1.38	1.99	2.77	2.07	2.49	2.33	3.56	3.56				
PLACE1005101	6.75	6.64	8.03	8.45	9.96	12.39	8.67	10.11	10.11		*	+	
PLACE1005102	5.88	7.51	8.49	11.05	10.78	12.60	9.73	9.59	9.59	*	+	+	
PLACE1005108	5.63	4.27	3.64	12.01	12.87	10.10	5.64	5.46	5.46	**	+		
PLACE1005110	6.84	3.16	2.29	5.61	4.42	2.27	2.47	3.96	3.96				
PLACE1005111	2.32	1.43	0.52	2.8	3.48	1.64	1.69	1.48	1.48				
PLACE1005123	20.53	8.57	10.06	12.54	14.07	10.45	7.24	8.30	8.3				
PLACE1005124	3.92	2.40	2.02	3.08	6.72	4.08	3.28	3.46	3.46				
PLACE1005128	10.6	9.42	9.74	12.9	15.61	15.03	14.09	17.89	17.89	**	+	**	+
PLACE1005130	4.63	4.42	3.58	6.21	6.12	6.60	2.90	3.62	3.62	**	+		
PLACE1005141	11.53	6.88	7.85	10.2	11.46	13.07	6.08	6.65	6.65				
PLACE1005146	2.66	2.45	2.31	3.79	4.23	2.90	1.91	2.35	2.35	*	+		
PLACE1005152	4.31	1.32	1.78	5.23	4.05	4.11	2.87	2.37	2.37				
PLACE1005157	3.17	1.71	2.58	3.61	2.97	3.04	1.83	2.24	2.24				
PLACE1005162	5.03	1.44	2.16	4.55	5.47	5.51	3.63	3.97	3.97				
PLACE1005170	1.73	0.31	0.62	1.61	1.26	1.41	1.34	1.72	1.72				
PLACE1005176	1.61	0.38	0.68	1.16	1.34	1.12	1.05	1.60	1.6				
PLACE1005181	0.5	0.24	0.53	1.19	0.87	2.59	0.77	1.26	1.26		*	+	
PLACE1005184	4.44	1.78	2.90	7.9	7.10	9.09	4.75	4.64	4.64	**	+		
PLACE1005186	6.95	2.41	3.82	3.37	3.80	2.87	3.22	3.68	3.68				
PLACE1005187	3.14	1.53	1.03	3.09	5.30	4.21	2.97	2.82	2.82				
PLACE1005189	5.93	2.53	2.32	3.58	5.81	4.44	5.57	5.74	5.74				
PLACE1005193	6.13	3.49	3.63	4.29	4.51	4.47	3.64	4.00	4				
PLACE1005200	4.37	1.39	2.33	2.59	3.60	1.69	2.29	2.95	2.95				
PLACE1005206	2.34	0.51	1.37	1.54	2.19	3.01	1.80	1.98	1.98				
PLACE1005216	1.38	0.71	1.11	2.26	2.41	2.76	2.43	3.73	3.73	**	+	**	+
PLACE1005223	4.29	2.34	2.64	6.04	7.76	7.97	4.06	6.10	6.1	**	+		
PLACE1005225	19.66	8.09	9.52	16.05	21.00	13.76	8.27	9.44	9.44				
PLACE1005232	8.02	4.04	2.69	6.94	10.56	7.61	5.96	6.58	6.58				
PLACE1005239	5.38	1.20	2.07	5.01	3.78	2.93	2.36	3.31	3.31				
PLACE1005243	5.32	3.76	4.72	5.19	5.09	5.33	3.34	5.82	5.82				
PLACE1005250	3.75	1.12	1.85	3.16	3.89	3.16	2.16	2.84	2.84				
PLACE1005261	2.07	0.70	1.90	2.25	2.05	1.77	2.13	1.93	1.93				
PLACE1005266	1.9	0.95	1.09	2.57	2.39	2.64	2.14	1.90	1.9	*	+		
PLACE1005271	5.66	2.63	3.94	8.71	9.11	8.37	4.71	5.02	5.02	**	+		
PLACE1005277	3.05	0.82	0.70	2.46	4.32	1.50	1.02	2.07	2.07				
PLACE1005287	6.59	3.30	3.94	10.35	15.42	7.57	8.69	8.45	8.45		*	+	
PLACE1005299	22.18	11.98	9.53	18.56	24.11	17.96	21.90	22.45	22.45				
PLACE1005305	5.96	2.44	4.52	8.17	10.96	9.42	8.88	11.22	11.22	*	+	**	+
PLACE1005307	3.74	1.42	2.86	4.85	5.32	3.53	2.69	4.11	4.11				
PLACE1005308	3.94	1.81	2.45	3.16	2.71	2.64	2.67	2.60	2.6				
PLACE1005313	1.8	1.22	2.93	1.89	0.89	2.76	1.70	1.69	1.69				
PLACE1005320	2.05	0.78	1.58	1.96	1.63	3.04	1.42	1.54	1.54				
PLACE1005327	3.57	2.45	2.12	2.64	6.29	3.81	4.41	6.45	6.45		*	+	
PLACE1005331	4	2.27	3.11	3.34	6.04	3.03	3.28	2.86	2.86				
PLACE1005335	9.31	5.05	4.18	8.68	7.24	5.98	5.53	6.95	6.95				
PLACE1005336	3.13	1.45	2.61	5.52	6.69	4.80	4.00	4.81	4.81	*	+	*	+
PLACE1005351	30.75	16.28	19.31	14.85	14.56	18.13	32.39	30.68	30.68				
PLACE1005366	3.38	2.74	2.56	10.21	9.37	10.62	9.15	9.50	9.5	**	+	**	+
PLACE1005373	4.26	1.58	2.70	3.39	2.69	4.82	2.63	3.29	3.29				

144/180

【0785】

【表314】

1765

1766

PLACE1005374	5	2.10	2.77	8.04	11.61	11.01	4.31	6.01	6.01	**	+		
PLACE1005383	8.86	3.18	3.37	5.63	6.03	4.19	5.25	6.23	6.23				
PLACE1005388	2.57	0.54	0.31	2.75	1.56	0.89	2.61	1.22	1.22				
PLACE1005409	3.48	3.06	2.63	7.59	8.06	6.25	3.31	4.02	4.02	*	+		
PLACE1005410	6.76	2.97	3.65	3.66	8.24	5.17	9.00	11.77	11.77		*	+	
PLACE1005426	4.46	1.72	1.45	2.27	1.48	1.00	3.43	3.54	3.54				
PLACE1005431	4.56	2.63	2.58	4.42	5.14	6.40	5.57	6.50	6.5		*	+	
PLACE1005453	3.55	1.77	2.09	4.33	4.49	5.14	1.74	3.20	3.2	*	+		
PLACE1005467	5.64	2.78	2.70	6.57	5.73	4.48	5.05	4.51	4.51				
PLACE1005471	3.36	0.50	1.20	3.42	3.09	2.65	2.30	3.64	3.64				
PLACE1005476	5.15	1.54	1.43	2.43	2.59	1.89	1.59	3.01	3.01				
PLACE1005477	2.24	1.35	1.27	5.66	7.05	5.00	4.23	7.05	7.05	**	+	*	+
PLACE1005488	1.93	1.39	1.29	1.24	1.52	1.24	1.31	1.75	1.75				
PLACE1005481	2.22	1.41	1.51	2.73	2.46	3.04	1.87	2.00	2	*	+		
PLACE1005494	1.24	0.38	0.90	0.8	0.90	0.66	0.80	1.98	1.98				
PLACE1005495	4.56	1.60	1.71	3.4	2.67	2.72	2.06	1.93	1.93				
PLACE1005497	8.06	4.83	3.69	4.42	2.88	4.07	9.50	10.40	10.4		*	+	
PLACE1005499	4.76	1.36	1.66	2.69	4.07	3.13	5.56	5.51	5.51				
PLACE1005500	2.69	0.87	1.10	2.75	3.41	2.24	1.89	4.02	4.02				
PLACE1005513	1.27	0.71	0.80	3.5	2.88	3.38	1.95	3.18	3.18	**	+	*	+
PLACE1005515	2.84	0.81	0.90	1.12	0.96	1.43	2.38	3.90	3.9				
PLACE1005519	7.14	2.92	5.14	2.37	3.46	3.11	2.55	3.35	3.35				
PLACE1005526	2.06	1.07	1.41	1.41	2.39	1.85	1.31	2.23	2.23				
PLACE1005528	6.82	2.99	3.77	7.7	10.09	11.05	4.64	5.96	5.96	*	+		
PLACE1005530	4.98	2.54	2.80	2.85	5.04	3.55	3.48	2.83	2.83				
PLACE1005536	4.27	3.13	1.98	6.1	4.77	1.67	4.10	3.87	3.87				
PLACE1005539	3	1.66	1.31	3.17	3.20	2.66	1.69	3.05	3.05				
PLACE1005543	2.3	1.25	1.18	4	3.96	4.38	3.55	3.32	3.32	**	+	**	+
PLACE1005544	6.06	3.23	2.89	3.81	4.11	4.35	4.12	5.12	5.12				
PLACE1005550	8.49	4.71	5.86	4.53	4.75	4.40	2.14	3.57	3.57				
PLACE1005554	1.55	0.76	0.94	1.77	1.45	1.38	2.99	1.56	1.56				
PLACE1005557	3.3	1.97	2.34	3.4	5.03	3.76	3.56	3.17	3.17				
PLACE1005563	1.99	2.09	0.76	1.69	2.10	1.89	2.11	1.69	1.69				
PLACE1005569	4.54	2.73	2.52	4.62	4.22	2.24	2.63	3.22	3.22				
PLACE1005574	1.43	0.92	0.87	2.29	2.41	2.10	0.45	0.99	0.99	**	+		
PLACE1005584	1.32	0.88	0.93	1.31	1.40	1.67	1.68	4.67	4.67				
PLACE1005590	2.53	3.81	2.63	3.18	2.75	3.39	4.08	5.93	5.93		*	+	
PLACE1005595	2.91	2.55	3.00	2.96	2.39	3.53	3.75	3.64	3.64		**	+	
PLACE1005601	2.77	1.99	2.02	2.52	2.79	3.50	2.97	3.86	3.86		*	+	
PLACE1005603	0.9	0.55	0.69	0.87	1.06	0.76	1.27	1.79	1.79		*	+	
PLACE1005604	4.18	2.56	1.82	4.89	4.83	6.27	2.39	1.93	1.93	*	+		
PLACE1005611	2.64	2.26	1.19	5.02	2.53	3.51	2.64	2.53	2.53				
PLACE1005622	2.15	1.96	1.00	2.49	2.91	2.25	1.48	2.00	2				
PLACE1005623	4.29	1.35	2.10	3.3	3.81	3.92	2.17	2.70	2.7				
PLACE1005630	6.26	3.63	2.27	4.66	6.06	5.41	4.45	5.87	5.87				
PLACE1005639	1.47	1.40	2.08	1.43	2.40	2.84	0.78	1.78	1.78				
PLACE1005646	5.91	4.51	5.24	4.63	5.74	5.46	4.51	5.47	5.47				
PLACE1005647	0.51	0.39	0.52	1.16	1.74	1.90	2.41	4.04	4.04	**	+	**	+
PLACE1005648	5.72	4.93	6.25	15.18	16.23	16.45	5.58	7.21	7.21	**	+		
PLACE1005653	3.3	1.90	0.82	3.94	5.11	4.03	2.85	2.07	2.07	*	+		
PLACE1005656	2.07	1.04	0.59	1.23	2.09	0.91	1.88	0.45	0.45				
PLACE1005659	4.14	1.56	2.46	2.97	4.17	2.22	1.44	2.31	2.31				
PLACE1005660	5.27	3.90	2.60	4.31	5.01	2.96	3.33	4.29	4.29				
PLACE1005664	4.13	4.07	4.07	5.57	5.47	4.07	5.14	6.25	6.25		**	+	
PLACE1005666	0.97	1.45	1.51	3.22	3.91	4.93	3.26	2.77	2.77	**	+	**	+
PLACE1005669	4.53	2.92	2.87	6.24	4.95	7.16	3.36	4.69	4.69	*	+		

145/180

1767

1768

PLACE1005683	2.11	2.05	2.13	4.34	3.23	4.41	1.89	2.15	2.15	**	+		
PLACE1005698	4.64	2.14	3.28	3.89	3.92	4.16	1.91	2.53	2.53				
PLACE1005708	25.78	13.70	10.51	13.88	16.18	11.27	14.00	14.43	14.43				
PLACE1005725	3.83	1.42	2.33	2.34	3.92	2.04	4.70	4.61	4.61		*	+	
PLACE1005727	8.48	2.80	3.97	5.4	4.41	4.96	2.49	2.57	2.57				
PLACE1005730	3.57	0.90	1.62	1.95	2.02	2.00	2.05	2.95	2.95				
PLACE1005734	4.39	2.36	2.88	8.34	10.28	9.63	5.13	7.81	7.81	**	+	+	+
PLACE1005739	2.31	1.03	1.11	1.47	1.17	1.64	2.72	2.15	2.15				
PLACE1005745	9.25	5.63	5.40	10.32	14.44	8.66	7.38	8.69	8.69				
PLACE1005752	4.63	2.11	0.91	2.57	2.97	2.88	2.25	2.86	2.86				
PLACE1005755	0.83	0.18	0.42	0.66	1.88	0.66	0.70	0.93	0.93				
PLACE1005756	14.63	7.31	9.39	22.2	25.42	27.72	29.92	35.68	35.68	**	+	**	+
PLACE1005760	7.89	3.72	4.80	10.59	12.05	10.96	9.43	9.92	9.92	*	+	+	+
PLACE1005763	3.86	1.70	3.26	6.59	6.36	6.88	4.43	4.28	4.28	**	+		
PLACE1005768	6.14	3.01	5.24	7.97	7.90	8.87	6.22	5.90	5.9	*	+		
PLACE1005771	7.62	3.12	5.03	7.4	7.32	9.76	6.04	6.48	6.48				
PLACE1005783	3.63	1.45	2.35	2.79	4.79	2.04	2.34	3.07	3.07				
PLACE1005799	6.45	3.16	3.38	5.32	4.64	3.49	5.15	5.23	5.23				
PLACE1005802	5.01	1.66	1.63	4.46	8.45	4.41	2.49	4.79	4.79				
PLACE1005803	11.48	4.59	6.77	9.23	10.65	9.39	6.33	8.91	8.91				
PLACE1005804	1.62	0.72	0.84	1.97	2.36	1.93	2.21	2.56	2.56	*	+	+	+
PLACE1005813	10.74	3.23	5.61	11.66	8.19	9.55	6.52	6.57	6.57				
PLACE1005815	5.12	2.48	3.85	7.34	9.35	11.87	4.89	5.17	5.17	*	+		
PLACE1005828	5.16	3.37	3.80	8.35	8.98	9.59	4.86	6.29	6.29	**	+		
PLACE1005833	3.06	1.35	1.59	18.69	21.23	11.91	28.00	30.88	30.88	**	+	**	+
PLACE1005834	1.93	0.65	0.55	4	6.43	2.66	1.50	2.50	2.5	*	+		
PLACE1005835	5.07	4.66	2.88	5.05	7.51	3.87	4.83	4.52	4.52				
PLACE1005836	3.75	1.63	2.11	2.62	6.42	3.23	2.73	2.06	2.06				
PLACE1005845	4.98	1.86	2.24	4.26	4.56	2.61	2.60	3.15	3.15				
PLACE1005850	4.23	2.74	2.58	5.55	4.59	5.10	2.95	3.19	3.19	*	+		
PLACE1005851	1.83	0.96	1.69	2.54	2.84	4.11	1.02	0.85	0.85	*	+		
PLACE1005856	4.08	1.29	7.53	4.1	2.89	3.39	1.78	2.05	2.05				
PLACE1005875	3.56	1.05	0.65	5.19	5.82	3.59	3.48	3.10	3.1				
PLACE1005876	4.08	3.91	2.72	2.79	2.82	2.10	2.04	2.27	2.27	*	-		
PLACE1005878	5.27	2.13	2.19	4.92	3.53	2.84	3.83	3.82	3.82				
PLACE1005880	3.44	0.96	1.32	2.14	2.64	2.46	2.97	4.34	4.34				
PLACE1005884	1.76	0.52	0.55	1.39	1.77	1.41	2.43	2.29	2.29		*	+	
PLACE1005890	2.04	0.70	0.65	1.41	1.86	1.52	1.88	2.21	2.21				
PLACE1005898	2.99	2.09	1.71	4.94	3.42	2.88	2.46	3.27	3.27				
PLACE1005913	5.71	2.57	3.76	7.83	8.39	8.51	3.79	4.62	4.62	*	+		
PLACE1005921	10.98	4.34	4.34	9.34	8.32	8.81	6.16	6.43	6.43				
PLACE1005923	57.96	26.97	25.39	4.09	4.25	2.49	3.95	3.48	3.48	*	-	*	-
PLACE1005925	2.51	0.91	2.14	3.11	3.71	2.82	1.93	2.80	2.8				
PLACE1005927	6.09	2.70	1.89	3.69	4.68	4.01	3.18	5.73	5.73				
PLACE1005932	1.82	0.71	0.41	1.33	1.76	1.08	1.15	1.54	1.54				
PLACE1005934	3.84	2.41	2.72	6.26	7.11	6.43	3.93	5.30	5.3	**	+	+	+
PLACE1005936	2.29	0.78	1.05	1.64	1.78	1.51	1.47	1.36	1.36				
PLACE1005939	6.69	4.35	3.92	5.44	5.46	3.73	16.40	25.30	25.3		**	+	
PLACE1005951	5.63	2.39	2.32	4.86	4.02	3.74	3.42	3.43	3.43				
PLACE1005953	2.92	1.24	0.96	2.51	1.78	2.30	1.15	2.55	2.55				
PLACE1005955	3.7	1.62	1.62	2.35	3.61	2.40	3.03	3.07	3.07				
PLACE1005966	3.38	0.65	1.08	1.86	2.32	2.12	2.02	2.80	2.8				
PLACE1005968	10.55	5.64	5.72	7.48	5.88	6.36	5.68	7.85	7.85				
PLACE1005975	10.44	5.19	8.44	12.95	14.63	15.64	8.66	16.92	16.92	*	+		
PLACE1005990	3.19	1.43	1.58	2.28	2.05	1.79	1.80	3.26	3.26				
PLACE1005997	64.81	36.05	40.42	54.4	53.64	53.12	27.58	33.55	33.55				

146/180

【0787】

【表316】

1769

1770

PLACE1006003	8.53	4.41	5.77	19.13	16.78	16.67	8.23	9.69	9.69	**	+		
PLACE1006003	6.88	5.62	5.05	3.42	5.00	5.45	4.05	7.43	7.43				
PLACE1006011	4.72	2.78	3.04	3.63	3.41	3.26	2.90	3.61	3.61				
PLACE1006017	4.17	1.57	1.37	3.12	3.78	3.87	3.13	4.29	4.29				
PLACE1006037	8.36	3.71	4.44	4.09	4.76	4.29	2.99	4.73	4.73				
PLACE1006040	13.34	8.65	10.10	9.09	7.82	11.18	9.13	10.46	10.46				
PLACE1006063	4.18	2.39	2.46	2.52	3.00	2.07	2.59	2.91	2.91				
PLACE1006071	3.1	2.05	2.07	1.68	2.75	3.43	1.83	2.76	2.76				
PLACE1006073	3.97	2.14	1.81	6.25	6.16	5.43	3.65	5.10	5.1	*	+		
PLACE1006074	4.44	2.36	2.42	6.36	6.76	5.83	2.98	4.13	4.13	*	+		
PLACE1006076	1.24	0.92	1.14	3.37	4.38	2.74	2.16	3.59	3.59	**	+	*	+
PLACE1006079	4.64	2.47	2.65	3.89	4.84	4.04	4.58	5.85	5.85				
PLACE1006093	1.06	0.90	1.72	1.34	1.63	0.86	2.10	2.38	2.38			*	+
PLACE1006116	2.79	1.95	1.97	2.66	2.53	2.69	3.38	3.33	3.33			*	+
PLACE1006119	2.59	2.94	2.87	5.28	4.68	6.57	3.23	3.84	3.84	**	+	*	+
PLACE1006129	2.82	1.25	0.50	2.84	2.73	3.10	3.07	1.53	1.53				
PLACE1006139	7.84	6.54	4.25	6.48	5.34	5.86	6.94	4.78	4.78				
PLACE1006143	2.36	1.84	1.60	4.6	3.86	4.22	1.68	3.18	3.18	**	+		
PLACE1006157	2.84	1.26	1.64	2.25	2.35	1.82	1.52	2.36	2.36				
PLACE1006159	1.74	1.38	1.27	2.48	3.25	2.76	3.72	4.61	4.61	**	+	**	+
PLACE1006164	0.77	0.31	0.34	1.19	1.94	1.20	1.01	0.99	0.99	*	+	*	+
PLACE1006167	6.97	5.82	7.53	6.63	9.38	9.10	8.80	7.88	7.88				
PLACE1006178	3.23	2.05	2.23	3.8	5.15	4.56	3.39	4.89	4.89	*	+	*	+
PLACE1006181	4.1	2.72	3.53	6.41	6.18	6.21	5.86	6.48	6.48	**	+	**	+
PLACE1006187	0.5	0.33	0.10	0.86	0.82	1.09	0.66	0.49	0.49	*	+		
PLACE1006195	3.24	1.23	1.17	2.67	2.87	2.14	2.62	1.30	1.3				
PLACE1006196	8.03	2.93	3.80	5.31	7.47	6.96	4.75	3.79	3.79				
PLACE1006197	7.57	3.83	6.49	6.35	7.27	5.99	3.44	4.86	4.86				
PLACE1006198	2.55	1.19	1.79	2.81	2.56	2.19	0.91	2.46	2.46				
PLACE1006205	0.84	0.89	1.05	0.57	0.49	1.57	0.74	1.36	1.36				
PLACE1006208	2.19	1.80	3.16	5	4.18	5.05	7.99	4.42	4.42	**	+		
PLACE1006211	24.46	16.18	17.64	12.62	6.69	13.24	6.25	5.01	5.01			**	-
PLACE1006219	3.37	2.25	3.36	4.14	6.29	3.89	6.74	5.53	5.53			**	+
PLACE1006223	1.64	1.06	2.11	4.34	4.03	4.26	2.35	1.45	1.45	**	+		
PLACE1006225	1.79	1.20	1.26	2	1.95	1.83	1.23	1.27	1.27				
PLACE1006236	1.44	1.01	1.87	3.01	4.09	1.96	1.59	2.02	2.02				
PLACE1006239	1.72	1.00	1.18	2.46	2.48	2.60	1.22	3.24	3.24	**	+		
PLACE1006245	3.4	2.04	2.24	3.29	3.77	3.95	2.39	2.28	2.28				
PLACE1006246	2.78	1.91	2.09	2.77	4.35	3.44	2.43	1.97	1.97				
PLACE1006248	1.93	1.11	1.30	3.09	2.89	3.27	2.22	1.63	1.63	**	+		
PLACE1006262	3.84	0.83	1.42	2.66	2.88	1.75	2.15	1.21	1.21				
PLACE1006269	3.04	1.04	0.97	1.76	1.81	1.87	1.28	1.88	1.88				
PLACE1006275	7.22	3.03	3.50	4.21	3.01	5.50	3.90	5.68	5.68				
PLACE1006277	2.96	1.17	2.13	3.73	2.55	2.62	1.86	2.11	2.11				
PLACE1006288	11.06	5.08	6.39	7.42	7.01	7.05	7.31	9.33	9.33				
PLACE1006290	2.57	0.88	2.08	1.76	1.36	2.22	3.49	3.56	3.56			*	+
PLACE1006298	4.88	1.93	2.02	5.27	5.25	6.12	2.74	3.06	3.06				
PLACE1006311	0.92	0.15	0.41	0.82	0.31	0.81	2.11	2.64	2.64			**	+
PLACE1006318	4.74	1.72	2.13	3.2	4.07	2.67	3.29	3.63	3.63				
PLACE1006325	9.29	2.77	3.97	3.52	11.00	5.12	4.02	5.45	5.45				
PLACE1006331	4.1	2.50	3.35	6.3	7.84	7.16	3.59	5.15	5.15	**	+		
PLACE1006335	4.07	1.71	1.37	3.4	2.75	3.34	2.92	1.94	1.94				
PLACE1006357	1.19	0.25	0.48	0.63	0.90	0.73	0.68	1.54	1.54				
PLACE1006360	5.46	2.85	2.81	3.84	5.35	5.87	2.18	2.76	2.76				
PLACE1006364	2.49	1.09	1.52	2.96	1.99	3.19	1.91	2.29	2.29				
PLACE1006365	0.49	0.34	1.09	1.19	1.22	2.88	0.94	1.48	1.48				

147/180

1772

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148/180

【表 3 1 8】

1774

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1776

PLACE1007092	13.8	11.82	5.85	6.03	7.76	3.70	4.55	4.38	4.38										
PLACE1007096	3.67	1.72	2.42	3.85	3.61	3.33	2.77	3.95	3.95										
PLACE1007097	2.22	0.99	0.99	1.67	2.32	2.35	3.22	1.09	1.09										
PLACE1007099	3.21	1.35	2.99	3.75	3.60	3.90	2.21	4.60	4.6										
PLACE1007105	3.27	1.47	1.70	2.02	1.66	2.46	1.10	2.81	2.81										
PLACE1007108	1.84	0.54	0.64	1.21	1.32	0.77	3.03	1.13	1.13										
PLACE1007111	1.12	0.75	0.77	2.41	0.87	1.64	1.17	1.43	1.43								*	+	
PLACE1007112	2.23	1.33	1.93	1.71	1.54	2.89	1.30	2.04	2.04										
PLACE1007130	1.72	0.36	0.26	1	1.71	0.63	0.85	1.29	1.29										
PLACE1007132	3.87	1.51	1.93	3.65	4.98	3.98	2.58	2.83	2.83										
PLACE1007140	2.78	1.67	1.49	5.51	4.02	1.94	1.59	4.61	4.61										
PLACE1007145	4.57	2.06	2.35	3.69	3.88	3.45	2.67	3.35	3.35										
PLACE1007169	7.86	3.91	6.07	4.6	3.97	4.34	4.66	5.06	5.06										
PLACE1007178	3.63	1.78	2.11	3.46	2.58	2.44	3.58	4.50	4.5										
PLACE1007190	1.52	0.85	1.18	1.02	0.96	1.35	1.62	1.51	1.51										
PLACE1007201	1.85	0.34	1.11	1.37	0.91	2.07	0.92	1.05	1.05										
PLACE1007202	18.75	9.75	12.22	19.49	17.57	13.05	23.70	22.24	22.24								*	+	
PLACE1007226	4.6	2.18	1.44	3.72	3.17	3.32	4.10	4.25	4.25										
PLACE1007238	4.59	1.78	4.87	4.05	4.43	2.63	3.54	2.85	2.85										
PLACE1007239	4.19	2.58	2.67	5.03	3.84	2.86	3.07	4.50	4.5										
PLACE1007242	3.6	1.20	1.84	1.27	2.10	2.41	1.99	2.58	2.58										
PLACE1007243	10.2	5.01	6.25	4.24	5.71	6.21	7.36	6.08	6.08										
PLACE1007247	3.28	2.10	1.67	14.75	8.63	15.61	4.03	8.60	8.6	**	+	*	+						
PLACE1007257	7.61	5.72	7.16	3.66	3.64	3.79	1.96	3.64	3.64	**	-	**	-						
PLACE1007274	4.38	2.42	3.36	7.38	8.79	6.79	3.07	4.64	4.64	**	+	*	+						
PLACE1007276	2.97	1.43	1.54	2.93	2.81	2.34	1.57	3.92	3.92										
PLACE1007282	8.6	4.51	8.76	10.51	12.35	10.29	22.66	27.14	27.14								**	+	
PLACE1007286	6	1.42	3.35	6.08	8.09	5.91	3.36	4.27	4.27										
PLACE1007296	5.96	3.96	4.56	9.09	9.08	8.48	6.51	8.92	8.92	**	+	*	+						
PLACE1007301	1.48	0.84	0.72	0.94	1.65	0.98	0.49	0.96	0.96										
PLACE1007314	7.72	5.09	4.39	7.99	9.50	9.98	8.19	8.10	8.1	*	+								
PLACE1007317	1.71	0.70	0.71	2.11	1.11	1.58	1.38	1.29	1.29										
PLACE1007329	1.19	1.05	0.73	3.19	2.34	1.79	1.73	2.65	2.65	*	+	*	+						
PLACE1007338	5.4	1.79	2.69	4.68	5.74	4.16	3.17	5.55	5.55										
PLACE1007343	2.46	2.38	1.97	2.04	2.30	2.39	2.65	5.91	5.91										
PLACE1007345	2.86	1.45	1.69	3.47	3.21	3.18	2.59	3.21	3.21	*	+								
PLACE1007346	5.8	4.00	4.67	8.73	7.57	8.39	4.92	8.73	8.73	**	+								
PLACE1007359	3.11	1.64	2.21	3.58	2.56	2.94	3.24	3.82	3.82										
PLACE1007367	9.92	5.57	8.83	12.43	19.19	16.79	8.33	10.26	10.26	**	+	*	+						
PLACE1007375	1.77	1.76	1.63	2.23	2.83	2.75	1.31	0.63	0.63	*	+	*	-						
PLACE1007377	4.63	2.52	2.53	3.52	3.56	1.75	2.11	3.18	3.18										
PLACE1007386	1.87	0.97	0.83	6.47	6.90	6.45	4.13	3.04	3.04	**	+	*	+						
PLACE1007392	2.72	3.07	3.82	2.83	2.94	3.03	2.89	3.43	3.43										
PLACE1007402	2.84	2.88	1.67	3.44	3.03	2.39	3.94	2.99	2.99										
PLACE1007409	0.93	0.91	1.34	1.36	1.18	1.53	1.35	1.51	1.51										
PLACE1007416	1.46	1.48	1.61	3.34	3.06	1.14	3.57	4.84	4.84	**	+	**	+						
PLACE1007420	9.86	15.04	12.94	15.11	14.93	22.60	15.85	14.30	14.3										
PLACE1007431	0.76	1.71	1.22	1.19	2.25	1.99	1.64	1.51	1.51										
PLACE1007450	4.02	1.67	1.64	5.21	4.82	5.24	2.44	2.49	2.49	*	+								
PLACE1007452	2.24	1.00	1.94	2.5	2.82	2.49	1.40	3.18	3.18										
PLACE1007454	10.17	5.34	5.33	10.29	13.21	14.02	9.07	11.14	11.14	*	+								
PLACE1007460	3.51	2.45	2.56	3.47	3.50	3.36	2.34	2.84	2.84										
PLACE1007478	1.85	1.34	0.98	2.14	2.65	2.21	0.62	1.89	1.89	*	+								
PLACE1007484	1.62	2.03	1.82	4.03	4.26	4.32	4.30	4.48	4.48	**	+	**	+						
PLACE1007488	2.83	1.13	1.39	2.08	1.66	1.77	1.01	1.64	1.64										
PLACE1007507	4.17	3.85	4.18	3.46	1.91	4.23	1.30	2.17	2.17	**	-	**	-						

1777

1778

PLACE1007511	1.09	1.11	0.68	1.33	1.45	0.75	0.90	1.48	1.48				
PLACE1007513	4.69	1.71	2.94	3.5	3.66	3.78	3.32	6.37	6.37				
PLACE1007524	6.92	2.48	2.90	3.93	4.08	2.82	1.80	1.66	1.66				
PLACE1007525	4.99	2.20	2.97	4.48	5.31	5.23	2.35	2.30	2.3				
PLACE1007537	3.67	3.75	2.72	3.67	3.58	4.70	2.62	4.19	4.19				
PLACE1007544	1.23	1.96	1.26	3.11	3.23	2.88	3.01	2.55	2.55	**	+	+	+
PLACE1007547	3.83	2.63	2.50	6.49	5.11	5.77	2.96	2.23	2.23	**	+		
PLACE1007557	3.78	2.86	3.01	6.18	5.42	6.26	3.20	3.81	3.81	**	+		
PLACE1007560	7.5	4.33	3.69	5.21	4.40	3.63	6.61	8.29	8.29				
PLACE1007565	1.39	0.57	0.51	1.55	0.69	1.08	1.27	0.93	0.93				
PLACE1007580	0.78	0.25	0.56	1.38	0.71	0.94	1.33	1.46	1.46		**	+	
PLACE1007583	1.68	1.21	1.36	3.07	1.74	2.51	1.23	2.34	2.34				
PLACE1007591	2.78	0.84	0.81	2.91	3.12	3.09	1.72	2.45	2.45				
PLACE1007598	4.1	2.36	3.10	8.03	7.01	9.10	4.75	4.36	4.36	**	+		
PLACE1007610	0.9	0.60	0.89	2.28	1.49	1.41	1.23	1.82	1.82	*	+	+	+
PLACE1007618	1.76	1.24	1.15	1.76	2.07	1.52	1.03	1.29	1.29				
PLACE1007631	2.86	1.26	1.24	2.73	3.31	2.18	1.97	2.67	2.67				
PLACE1007634	6.13	3.63	3.43	16.1	18.88	18.33	14.85	19.91	19.91	**	+	**	+
PLACE1007632	4.92	2.23	3.27	3.4	3.01	3.01	4.94	4.29	4.29				
PLACE1007635	3.04	0.96	2.65	2.16	2.56	2.69	1.76	2.94	2.94				
PLACE1007645	4.04	1.20	2.15	4.72	5.27	5.01	4.78	4.87	4.87	*	+	+	+
PLACE1007649	1.28	0.79	0.67	1.29	1.36	2.38	1.28	2.15	2.15				
PLACE1007652	4.23	1.93	2.69	6.75	3.97	6.88	2.94	4.41	4.41				
PLACE1007669	6.2	1.80	2.99	5.47	6.53	4.51	3.57	2.86	2.86				
PLACE1007677	4.22	1.89	1.71	6.84	8.75	7.28	3.90	4.46	4.46	**	+		
PLACE1007688	5.22	1.69	2.55	2.63	3.33	2.71	2.38	2.43	2.43				
PLACE1007690	3.97	2.16	3.39	4.09	4.66	3.97	3.53	4.50	4.5				
PLACE1007697	1.72	0.75	0.98	1.08	0.70	0.98	1.28	0.95	0.95				
PLACE1007702	1.76	0.86	1.32	1.85	1.37	3.00	2.01	1.95	1.95				
PLACE1007705	2.4	0.53	1.89	1.45	2.19	2.67	2.64	2.34	2.34				
PLACE1007706	2.8	1.14	1.84	2.88	2.31	2.20	2.45	2.27	2.27				
PLACE1007725	3.27	2.02	1.52	3.44	3.01	2.26	1.89	1.39	1.39				
PLACE1007729	3.75	0.91	0.48	1.28	1.88	1.09	1.35	1.46	1.46				
PLACE1007730	4.12	1.63	2.33	3.92	2.43	2.55	1.94	4.18	4.18				
PLACE1007737	4.58	2.53	1.58	4.31	5.53	6.14	3.60	3.45	3.45				
PLACE1007743	1.47	0.73	0.61	2.7	2.78	2.53	1.94	2.71	2.71	**	+	+	+
PLACE1007746	3.82	1.81	2.10	5.73	3.58	6.69	6.74	9.08	9.08	**	+		
PLACE1007753	2.19	1.29	1.71	1.02	1.20	1.89	1.49	1.55	1.55				
PLACE1007769	0.98	0.53	0.69	1.58	1.14	1.77	1.01	1.04	1.04	*	+		
PLACE1007780	4.5	2.26	1.99	3.89	4.09	2.46	2.36	2.20	2.2				
PLACE1007791	5.12	2.18	2.04	3.75	4.60	3.26	2.31	3.66	3.66				
PLACE1007807	2.35	0.20	1.17	3.74	3.71	3.65	3.45	3.14	3.14	*	+	+	+
PLACE1007810	1.24	0.07	0.47	1.06	0.82	1.32	1.17	1.10	1.1				
PLACE1007814	5.26	2.80	2.95	4.73	4.47	4.22	4.12	5.14	5.14				
PLACE1007828	1.64	1.27	1.04	1.35	1.67	1.95	1.48	2.89	2.89				
PLACE1007829	6.87	2.06	4.61	11.59	10.29	14.12	4.54	6.24	6.24	*	+		
PLACE1007841	2.09	0.69	0.83	1.22	2.33	3.41	1.28	2.06	2.06				
PLACE1007842	2.47	1.09	2.35	2.63	2.75	2.08	1.49	2.36	2.36				
PLACE1007843	1.12	0.63	0.54	0.94	1.58	0.91	0.72	0.88	0.88				
PLACE1007845	3.75	1.17	2.10	1.73	2.80	1.91	2.13	2.52	2.52				
PLACE1007846	4.22	1.07	1.41	5.34	3.41	3.57	3.77	4.73	4.73				
PLACE1007848	1.96	0.65	0.52	1.52	2.21	2.88	2.45	2.38	2.38		*	+	
PLACE1007851	2.98	0.96	2.30	2.1	4.32	3.04	2.76	3.44	3.44				
PLACE1007858	1.43	0.68	1.60	6.59	6.03	5.72	3.52	4.91	4.91	**	+	**	+
PLACE1007866	30.58	17.58	17.98	11.58	9.80	8.70	10.80	10.41	10.41				
PLACE1007871	22.99	9.51	12.34	18.55	16.06	17.86	13.04	17.83	17.83				

151/180

【0792】

【表321】

1779

1780

PLACE1007877	4.54	1.36	1.17	4.16	4.25	2.80	3.39	3.32	3.32										
PLACE1007878	4.4	2.07	2.29	2.41	2.70	2.37	3.13	5.04	5.04										
PLACE1007881	1.27	0.74	0.75	0.94	1.76	0.67	0.87	1.11	1.11										
PLACE1007885	1.23	1.17	1.11	1.97	2.06	1.97	2.46	3.25	3.25	**	+	**	+						
PLACE1007897	2.56	0.68	1.11	1.75	1.79	1.30	1.00	2.88	2.88										
PLACE1007908	7.68	3.04	3.27	4.73	4.71	5.04	4.39	4.18	4.18										
PLACE1007922	1.4	0.69	0.89	1.56	0.63	1.43	1.13	0.93	0.93										
PLACE1007946	4.36	3.22	3.12	4.56	4.09	3.11	2.97	3.28	3.28										
PLACE1007950	5.15	1.51	1.60	3.7	3.21	2.35	3.25	8.99	8.99										
PLACE1007954	3.66	2.15	2.27	2.4	2.26	2.19	2.79	1.92	1.92										
PLACE1007955	4.71	1.37	1.67	2.61	3.53	2.54	2.49	4.46	4.46										
PLACE1007956	4.42	1.04	2.64	3.61	3.50	3.32	2.21	3.84	3.84										
PLACE1007958	1.93	0.27	1.12	1.34	1.94	1.66	1.60	1.84	1.84										
PLACE1007965	2.55	1.76	1.99	2.32	2.51	3.02	1.19	2.52	2.52										
PLACE1007969	6.03	2.86	2.43	4.73	5.79	6.79	4.72	3.77	3.77										
PLACE1007971	3.53	1.27	2.02	3.82	4.31	3.71	3.34	3.31	3.31										
PLACE1007990	2.84	1.35	1.80	4.92	3.19	2.61	2.45	2.53	2.53										
PLACE1008000	1.73	0.77	0.35	3.42	1.14	0.76	1.28	1.93	1.93										
PLACE1008002	0.38	0.09	0.23	1.64	0.83	0.73	1.52	1.90	1.9	*	+	**	+						
PLACE1008037	0.98	0.19	0.99	1.13	1.05	1.34	1.22	1.68	1.68										
PLACE1008044	4.87	3.62	2.89	3.52	3.76	3.71	2.56	3.52	3.52										
PLACE1008045	1.81	1.03	1.31	1.51	1.59	1.22	1.49	2.12	2.12										
PLACE1008080	4.1	3.05	2.36	3.11	3.91	2.99	2.39	3.89	3.89										
PLACE1008092	2.02	1.71	1.46	1.1	0.88	0.81	1.07	2.15	2.15	*	-								
PLACE1008095	2.93	1.27	1.19	2.55	1.83	2.32	1.34	3.34	3.34										
PLACE1008105	2.48	0.98	1.47	2.27	0.97	1.49	2.91	5.54	5.54					*	+				
PLACE1008107	6.58	3.57	3.85	1.29	1.19	1.39	4.33	5.78	5.78	*	-								
PLACE1008111	2.46	1.02	2.41	3.33	2.35	3.47	2.96	3.00	3										
PLACE1008113	25.85	13.24	14.36	22.14	19.88	22.12	9.93	8.27	8.27										
PLACE1008122	1.07	0.36	1.70	1.64	1.18	1.29	1.04	1.29	1.29										
PLACE1008129	1.31	1.01	1.72	3.06	3.91	4.22	1.89	1.53	1.53	**	+								
PLACE1008132	2.89	1.43	1.69	4.85	4.46	4.06	3.75	2.77	2.77	**	+								
PLACE1008137	3.98	1.85	1.77	2.91	2.34	1.96	2.43	2.78	2.78										
PLACE1008174	10.37	5.11	6.06	7.46	7.08	5.83	3.58	4.68	4.68										
PLACE1008177	5.22	2.35	2.42	4.78	5.45	4.55	2.08	2.73	2.73										
PLACE1008181	0.6	0.35	0.59	2.1	1.63	0.83	0.78	0.73	0.73					*	+				
PLACE1008195	4.21	3.69	4.41	3.34	3.31	4.29	3.54	5.03	5.03										
PLACE1008198	0.92	1.28	1.62	1.49	2.09	2.17	1.39	2.32	2.32										
PLACE1008201	1.66	0.51	1.49	2.83	2.14	2.43	2.07	1.72	1.72	*	+								
PLACE1008209	5.39	4.27	2.17	7.66	7.83	6.93	6.08	4.07	4.07	*	+								
PLACE1008226	3.09	1.71	1.62	2.88	3.33	2.83	2.61	2.73	2.73										
PLACE1008227	3.17	1.23	2.12	4.9	4.87	5.42	2.16	2.72	2.72	**	+								
PLACE1008231	2.12	0.50	0.70	1.87	1.47	1.28	1.21	0.99	0.99										
PLACE1008238	3.15	3.76	3.38	3.65	4.20	4.83	4.89	4.62	4.62			**	+						
PLACE1008244	1.2	0.39	0.55	1.23	1.76	1.25	0.99	1.37	1.37										
PLACE1008249	2.18	0.27	0.99	2.07	1.33	1.47	0.79	1.16	1.16										
PLACE1008266	3.92	3.58	3.56	7.01	10.06	9.34	6.60	6.76	6.76	**	+	**	+						
PLACE1008273	2.91	1.72	1.49	4.31	6.22	5.51	5.60	5.70	5.7	**	+	**	+						
PLACE1008275	1.29	0.61	1.24	2.1	1.76	1.18	1.34	0.60	0.6										
PLACE1008280	2.51	1.09	1.40	1.61	2.66	1.55	2.19	1.36	1.36										
PLACE1008282	6.02	2.61	4.50	6.93	7.87	7.46	6.98	6.73	6.73	*	+								
PLACE1008297	1.93	0.37	0.80	1.67	1.93	1.78	1.68	1.21	1.21										
PLACE1008303	2.86	2.08	2.50	1.98	2.20	2.57	2.27	1.77	1.77										
PLACE1008309	1	0.36	0.94	1.24	0.57	1.29	1.65	0.87	0.87										
PLACE1008315	12.99	6.08	6.62	5.63	6.15	4.10	5.15	5.08	5.08										
PLACE1008329	5.4	1.64	1.66	3.46	3.09	2.21	1.61	3.12	3.12										

152/180

【0793】

【表322】

1782

PLACE1008330	3.99	1.02	3.12	3.69	2.72	3.55	2.59	3.30	3.3										
PLACE1008331	3.5	1.58	2.61	2.43	4.87	4.55	2.21	5.77	5.77										
PLACE1008331	3.59	1.91	2.57	5.18	5.19	5.56	3.81	3.50	3.5	**	+								
PLACE1008354	3.92	0.69	2.72	2.64	2.56	2.29	2.42	2.95	2.95										
PLACE1008359	1.48	0.76	0.90	2.22	1.26	2.34	1.68	2.46	2.46								*	+	
PLACE1008368	4.18	1.66	2.15	9.15	7.54	8.92	6.11	7.44	7.44	**	+	**							
PLACE1008369	2.77	0.73	1.19	2.41	7.30	3.35	1.02	1.60	1.6										
PLACE1008392	2.13	0.98	1.09	1.58	3.18	1.77	1.88	2.10	2.1										
PLACE1008394	26.4	13.24	13.94	17.36	15.53	22.06	16.70	19.87	19.87										
PLACE1008398	7.2	3.44	10.45	4.58	8.83	4.91	2.86	4.01	4.01										
PLACE1008401	3.08	0.75	1.07	1.76	1.56	2.79	1.84	3.10	3.1										
PLACE1008402	6.01	1.01	4.48	2.49	3.09	3.48	2.05	3.35	3.35										
PLACE1008405	25.84	13.96	18.38	38.51	28.28	49.12	27.91	33.39	33.39								*	+	
PLACE1008409	16.67	9.53	11.29	12.69	10.07	15.56	12.51	11.76	11.76										
PLACE1008420	5.7	4.00	2.86	5.32	4.44	3.71	4.42	4.23	4.23										
PLACE1008424	3.57	2.25	1.23	2.09	2.46	2.00	2.48	2.25	2.25										
PLACE1008426	4.1	1.19	2.55	2.53	2.76	1.73	1.42	1.69	1.69										
PLACE1008429	1.34	0.85	1.46	2	3.50	1.65	1.93	1.52	1.52										
PLACE1008430	1.82	0.58	0.88	2.02	1.64	0.56	0.86	2.26	2.26										
PLACE1008437	2.06	0.49	1.54	1.53	1.27	1.54	1.33	2.88	2.88										
PLACE1008453	3.99	2.14	2.45	2.78	2.86	2.41	2.29	5.19	5.19										
PLACE1008454	4.67	3.03	4.69	8.04	6.50	8.39	3.85	5.65	5.65	*	+								
PLACE1008455	6.35	2.17	1.87	10.14	10.23	5.77	6.05	5.82	5.82										
PLACE1008457	9.43	3.52	3.32	5.83	7.73	6.63	5.24	7.01	7.01										
PLACE1008465	2.14	1.13	1.61	1.53	3.02	1.33	2.20	2.70	2.7										
PLACE1008469	12.37	7.23	7.87	8.96	9.09	12.38	13.17	10.93	10.93										
PLACE1008488	1.94	0.92	1.25	0.9	1.06	1.44	1.44	0.95	0.95										
PLACE1008519	3.83	1.77	1.73	2.4	1.77	1.88	2.77	1.49	1.49										
PLACE1008524	3.06	0.85	1.87	3.33	2.40	3.53	2.10	1.92	1.92										
PLACE1008531	3.02	1.05	2.48	2.83	2.67	2.71	2.79	2.45	2.45										
PLACE1008532	1.95	1.34	1.62	3.81	2.59	2.68	2.83	3.90	3.9	*	+	**	+						
PLACE1008533	6.08	2.16	3.15	4.18	5.64	3.25	3.67	5.24	5.24										
PLACE1008542	3.98	1.49	1.76	4.67	6.17	4.59	3.86	6.21	6.21	*	+								
PLACE1008549	2.51	1.53	0.88	1.7	2.51	1.76	1.36	1.66	1.66										
PLACE1008560	1.85	0.72	0.75	0.85	0.84	0.96	2.24	1.41	1.41										
PLACE1008567	2.83	1.62	2.07	2.6	2.14	2.90	2.18	3.74	3.74										
PLACE1008568	1.44	0.85	1.22	4.02	2.55	4.05	2.96	3.07	3.07	*	+	**	+						
PLACE1008569	6.68	1.97	2.63	4.52	4.62	4.72	3.58	5.21	5.21										
PLACE1008584	2.8	0.91	1.34	2.88	1.91	1.76	1.37	1.81	1.81										
PLACE1008585	6.05	2.08	1.87	5.97	6.88	5.16	6.30	6.66	6.66										
PLACE1008603	2.79	0.92	1.64	1.88	1.63	1.46	1.46	2.30	2.3										
PLACE1008621	2.19	0.44	1.30	1.02	1.47	0.69	1.47	2.18	2.18										
PLACE1008625	0.9	0.37	0.80	0	0.79	0.63	1.51	1.36	1.36								*	+	
PLACE1008626	1.01	0.36	0.40	1.03	0.59	0.80	0.60	2.30	2.3										
PLACE1008627	3.31	1.35	1.85	3.04	2.64	2.27	2.82	2.83	2.83										
PLACE1008629	4.46	2.86	3.88	4.95	3.86	3.87	2.66	4.45	4.45										
PLACE1008630	6.49	3.28	4.20	4.75	4.80	4.92	3.62	3.61	3.61										
PLACE1008643	3.94	1.90	2.23	4.63	3.91	2.95	3.01	3.94	3.94										
PLACE1008650	1.04	0.28	0.89	1.14	0.65	0.67	0.98	2.48	2.48										
PLACE1008657	2.91	1.23	0.78	2.05	1.78	1.50	2.02	2.34	2.34										
PLACE1008664	2.55	1.44	2.26	1.59	2.27	1.94	2.31	1.74	1.74										
PLACE1008693	3.83	1.61	1.78	3.36	3.20	3.43	2.93	2.63	2.63										
PLACE1008696	1.57	0.88	1.00	2.25	2.32	2.53	1.82	1.87	1.87	**	+	*	+						
PLACE1008715	1.2	1.08	0.38	2.73	1.58	1.58	1.30	1.20	1.2										
PLACE1008716	2.62	1.18	1.33	2.82	3.53	2.15	2.83	1.88	1.88										
PLACE1008722	8.81	3.15	4.14	9.07	11.88	9.16	5.01	7.77	7.77										

153/180

1784

PLACE1008738	1.83	2.28	2.00	1.8	1.24	1.00	1.36	3.09	3.09										
PLACE1008742	4.02	1.70	1.54	4.3	5.17	3.46	2.80	3.04	3.04										
PLACE1008744	1.17	0.49	0.67	1.04	1.21	1.19	1.00	1.69	1.69										
PLACE1008748	1.18	0.53	1.02	1.35	1.38	1.66	1.55	1.10	1.1										
PLACE1008757	0.57	0.66	1.64	0.86	1.31	1.19	0.28	1.35	1.35										
PLACE1008766	5.2	1.84	3.38	5.73	6.06	11.79	4.24	3.09	3.09										
PLACE1008785	3.43	1.55	1.67	3.73	3.48	3.51	2.86	2.40	2.4										
PLACE1008790	4.68	2.15	2.15	5.43	4.49	3.61	3.28	3.45	3.45										
PLACE1008791	6.35	0.62	2.86	2.36	3.47	2.69	1.71	2.65	2.65										
PLACE1008807	0.99	1.20	1.36	0.98	1.48	1.58	0.90	2.29	2.29										
PLACE1008808	2.02	1.19	1.16	1.26	1.76	1.00	2.24	1.72	1.72										
PLACE1008813	0.94	0.76	1.96	0.73	1.40	0.71	0.81	2.94	2.94										
PLACE1008836	3.35	2.03	2.82	3.36	3.83	3.93	1.76	4.97	4.97										
PLACE1008851	6.37	2.37	2.20	3.21	3.73	4.45	1.84	2.02	2.02										
PLACE1008854	1.01	0.67	0.67	0.73	1.08	1.01	0.89	0.70	0.7										
PLACE1008864	5.23	2.45	2.26	6.92	5.99	5.19	3.11	3.68	3.68										
PLACE1008867	1.96	1.35	1.26	5.74	4.65	5.92	4.30	4.51	4.51	**	+								
PLACE1008876	51.43	26.54	27.05	38	43.35	42.72	24.30	22.52	22.52										
PLACE1008887	1.78	0.54	1.07	2.31	2.39	2.93	1.78	2.61	2.61	*	+								
PLACE1008902	1.97	0.82	0.85	1.66	1.42	3.56	1.02	2.90	2.9										
PLACE1008911	6.01	5.11	5.63	8.6	8.99	8.79	6.07	6.33	6.33	**	+								
PLACE1008917	3.34	2.37	2.25	2.63	3.74	3.27	2.99	3.43	3.43										
PLACE1008929	1.57	0.52	0.53	1.3	2.33	1.36	0.77	1.37	1.37										
PLACE1008925	1.43	1.01	0.48	2.16	1.60	0.85	1.24	0.93	0.93										
PLACE1008938	8.48	4.04	4.74	5.59	5.27	6.20	2.97	5.51	5.51										
PLACE1008934	2.73	1.83	1.68	2.96	2.07	1.68	2.13	1.92	1.92										
PLACE1008941	2.12	4.29	2.29	2.81	3.70	3.18	1.74	1.69	1.69	*	+	**							
PLACE1008947	5.3	4.86	3.97	6.01	5.96	5.46	4.91	5.47	5.47										
PLACE1008954	2.32	1.08	1.90	4.47	4.44	4.99	1.56	2.13	2.13	**	+								
PLACE1008985	1.06	1.41	1.57	2.31	2.24	1.90	1.22	3.49	3.49	*	+								
PLACE1008994	1.26	0.32	0.61	1.19	2.34	0.75	0.51	0.61	0.61										
PLACE1008920	2.03	0.83	0.79	1.36	0.98	0.99	0.91	1.17	1.17										
PLACE1009027	2.42	0.29	0.98	17.03	20.58	24.13	13.27	17.48	17.48	**	+	**							
PLACE1009039	0.66	0.39	0.60	0.97	0.77	0.82	0.81	1.68	1.68	*	+	*							
PLACE1009045	1.25	0.20	1.18	0.92	1.61	1.30	3.10	3.19	3.19			**	+						
PLACE1009048	0.29	0.37	0.55	0.51	0.66	0.96	1.73	0.67	0.67										
PLACE1009050	0.48	(0.04)	0.53	1.13	0.72	1.09	0.42	0.86	0.86	*	+								
PLACE1009060	3.31	1.27	1.72	4.36	1.92	4.74	2.50	4.91	4.91										
PLACE1009067	4.9	1.27	1.78	2.92	1.97	2.26	4.68	4.77	4.77										
PLACE1009071	5.93	4.97	3.58	6.84	4.81	6.47	5.46	4.55	4.55										
PLACE1009090	3.14	0.90	2.12	3.01	2.91	3.24	2.46	1.95	1.95										
PLACE1009091	4.11	1.05	1.26	1.69	2.73	1.26	0.58	1.98	1.98										
PLACE1009094	2.34	2.30	1.26	2.48	1.83	1.50	3.22	2.13	2.13										
PLACE1009099	4.71	2.33	2.35	5.94	5.89	8.61	3.69	5.79	5.79	*	+								
PLACE1009110	1.06	1.13	0.63	4.86	1.21	3.08	2.60	2.41	2.41			**	+						
PLACE1009111	1.61	0.55	0.64	2.6	1.26	1.76	1.01	2.06	2.06										
PLACE1009113	5.16	1.93	2.40	3.84	4.61	2.47	1.71	4.56	4.56										
PLACE1009130	2.4	1.03	1.11	1.45	1.93	2.94	1.60	1.65	1.65										
PLACE1009150	1.73	0.66	1.55	2.16	2.01	2.30	1.65	1.47	1.47										
PLACE1009155	3.13	2.31	1.89	4.69	5.44	5.47	2.82	2.95	2.95	**	+								
PLACE1009158	3.54	1.36	1.91	2.88	2.25	2.53	2.92	2.13	2.13										
PLACE1009166	2.58	1.73	2.09	2.03	2.15	2.17	2.50	2.39	2.39										
PLACE1009172	2.84	0.78	1.90	4.25	3.46	3.67	2.50	4.14	4.14	*	+								
PLACE1009174	3.1	1.74	1.40	4.47	5.90	4.15	2.46	2.47	2.47	*	+								
PLACE1009183	6.02	1.51	2.01	3.08	4.98	3.03	2.18	2.53	2.53										
PLACE1009186	3.59	0.98	1.37	2.08	2.13	0.57	1.69	3.99	3.99										

154/180

【0795】

【表 3 2 4】

1785

1786

FLACE1009190	2.12	1.27	2.18	1.35	2.00	2.47	0.78	2.21	2.21								
FLACE1009196	1.64	0.69	1.48	2.04	2.57	3.98	1.85	1.52	1.52								
FLACE1009200	4.32	1.09	2.61	4.48	5.35	4.97	2.74	2.68	2.68								
FLACE1009217	2.54	0.82	0.83	0.92	1.24	1.76	2.27	2.78	2.78								
FLACE1009230	3.29	1.25	2.57	3.85	3.86	4.23	1.77	4.02	4.02								
FLACE1009236	3.68	1.44	1.56	2.57	2.82	2.63	1.54	2.09	2.09								
FLACE1009246	9.73	3.62	4.17	6.98	7.72	5.06	6.33	5.96	5.96								
FLACE1009265	21.04	8.85	7.61	12.85	14.86	12.34	4.96	7.60	7.6								
FLACE1009279	1.84	0.86	0.79	1.58	1.52	1.53	1.15	1.01	1.01								
FLACE1009288	3.7	2.72	2.61	7.54	8.77	8.06	7.00	9.82	9.82	**		+	**				
FLACE1009309	8.08	4.61	4.25	6.42	4.02	4.44	5.48	7.05	7.05								
FLACE1009319	2.03	1.05	1.47	2.87	1.77	3.10	1.90	2.70	2.7								
FLACE1009328	1.59	0.99	1.42	4.54	4.75	5.66	3.66	4.23	4.23	**		+	**				
FLACE1009335	1.22	0.54	0.61	2.18	1.74	1.92	1.46	0.54	0.54	**		+	**				
FLACE1009338	3.48	1.35	1.84	5.85	6.71	4.36	2.31	2.98	2.98	*							
FLACE1009344	3.01	1.13	2.79	1.83	3.29	2.00	2.97	2.70	2.7								
FLACE1009355	1.86	0.75	0.42	1.64	1.55	1.14	2.65	5.34	5.34								
FLACE1009366	2.14	1.43	1.26	1.31	1.41	1.74	1.22	2.07	2.07								
FLACE1009375	1.44	0.73	1.31	0.98	2.28	1.80	1.47	2.25	2.25								
FLACE1009388	1.69	1.27	1.19	3.96	2.82	3.05	1.65	2.75	2.75	**		+	**				
FLACE1009398	6.96	2.57	3.77	9	5.66	6.33	4.19	4.18	4.18								
FLACE1009404	4.11	2.25	3.40	3.14	5.18	4.09	2.94	3.62	3.62								
FLACE1009410	1.58	0.66	0.54	0.77	1.47	0.75	1.04	1.03	1.03								
FLACE1009417	1.85	0.80	1.11	2.36	1.87	0.83	3.31	3.04	3.04								
FLACE1009424	10.71	5.65	7.84	8.47	7.50	6.48	8.06	10.17	10.17								
FLACE1009434	3.29	1.53	1.47	2.38	1.85	1.49	1.58	1.71	1.71								
FLACE1009443	2.96	1.10	1.13	1.36	1.62	1.85	0.98	1.60	1.6								
FLACE1009444	3.55	2.71	1.84	4.89	4.13	5.32	3.26	4.47	4.47	*		+	*				
FLACE1009459	5.23	2.29	2.82	3.92	3.20	3.43	3.08	4.21	4.21								
FLACE1009460	0.43	0.37	0.33	0.44	0.37	1.88	0.42	0.69	0.69								
FLACE1009468	5.92	2.35	2.32	5.54	2.65	2.84	4.15	2.97	2.97								
FLACE1009476	2.6	0.89	1.54	2.02	2.17	1.83	1.69	2.92	2.92								
FLACE1009477	3.84	1.44	1.65	4.37	2.97	3.00	2.09	2.93	2.93								
FLACE1009483	2.08	0.70	1.33	2.12	1.22	1.24	0.82	2.09	2.09								
FLACE1009502	0.95	0.44	0.76	0.93	0.90	0.72	0.86	1.97	1.97								
FLACE1009524	2.21	0.79	1.36	1.49	1.81	1.15	1.58	2.15	2.15								
FLACE1009527	1.81	1.71	1.43	2.21	1.72	1.29	1.43	1.91	1.91								
FLACE1009531	5.24	3.01	2.51	5.69	3.69	5.37	6.78	6.24	6.24								
FLACE1009539	1.5	0.44	0.55	2.44	1.80	1.96	2.38	1.44	1.44	*							
FLACE1009539	3.39	1.25	2.38	2.92	3.40	3.47	2.40	3.54	3.54								
FLACE1009540	6	3.37	5.39	4.83	4.41	4.48	4.99	6.08	6.08								
FLACE1009543	2.35	1.42	1.51	1.82	1.71	1.38	1.98	2.97	2.97								
FLACE1009546	1.47	0.53	0.69	0.94	1.26	0.62	1.78	0.85	0.85								
FLACE1009556	1.35	0.95	1.07	1.35	2.21	0.98	2.07	2.20	2.2					**	+		
FLACE1009569	2.13	1.30	1.80	2.87	2.82	3.16	1.76	2.07	2.07	*		+	*				
FLACE1009571	2.72	1.88	1.50	2.08	1.82	2.50	1.30	1.73	1.73								
FLACE1009573	8.32	4.58	4.70	4.98	4.59	3.46	3.53	2.68	2.68								
FLACE1009576	3.44	1.43	2.32	5.25	3.67	4.91	3.85	4.08	4.08	**		+	**				
FLACE1009580	2.8	1.13	1.78	3.7	3.05	2.82	3.23	3.94	3.94					*	+		
FLACE1009581	2.06	1.05	0.67	2.39	2.40	2.16	4.59	3.83	3.83					**	+		
FLACE1009587	1.75	1.08	0.69	1.11	1.37	1.41	1.48	2.01	2.01								
FLACE1009593	2.92	1.61	2.66	2.04	2.95	2.35	2.20	2.52	2.52								
FLACE1009595	4.18	2.88	2.05	6.18	7.63	5.15	3.71	4.73	4.73	*		+	*				
FLACE1009596	1.65	1.09	0.96	1.87	2.16	2.45	1.64	1.48	1.48	*		+	*				
FLACE1009600	6.27	3.95	2.87	7.95	6.43	4.31	4.05	5.38	5.38								
FLACE1009604	2.52	0.69	0.99	3.24	2.45	1.67	2.85	2.11	2.11								

1787

1788

PLACE1002607	3.67	1.38	1.49	4.1	6.22	4.84	3.49	3.02	3.02	*	+		
PLACE1002613	3.3	1.40	2.05	3.5	4.21	3.44	2.36	3.25	3.25				
PLACE1002621	2.39	1.87	2.42	5.45	5.01	5.43	4.10	6.00	6	**	+	**	+
PLACE1002623	1.78	0.78	1.73	2.06	1.60	1.99	2.28	4.60	4.6		*		+
PLACE1002624	0.78	1.54	0.90	2.28	3.24	1.75	2.30	1.54	1.54				
PLACE1002637	1.33	0.77	0.84	3.69	2.89	3.73	3.44	3.33	3.33	**	+	**	+
PLACE1002639	2.08	0.08	0.65	2.19	1.89	1.62	1.82	1.57	1.57				
PLACE1002654	2.53	0.76	1.11	2.83	1.57	1.39	1.86	2.31	2.31				
PLACE1002659	5.89	3.14	3.71	3.85	5.36	4.32	4.03	5.80	5.8				
PLACE1002665	1.27	1.04	0.92	2.92	2.14	2.54	0.79	2.03	2.03	**	+		
PLACE1002669	3.5	3.60	3.11	3.69	5.37	3.54	3.97	4.99	4.99		*	+	
PLACE1002670	2.16	1.80	1.32	3.29	1.88	3.37	1.92	2.64	2.64				
PLACE1002788	2.48	1.90	1.93	4.13	3.31	5.20	2.14	3.90	3.9	*	+		
PLACE1002721	3.15	2.27	2.41	1.67	3.17	2.28	7.20	2.48	2.48				
PLACE1002731	3.26	1.56	1.59	2.49	3.83	1.81	1.89	2.37	2.37				
PLACE1002735	2.96	1.31	2.04	2.52	2.63	2.49	2.46	2.74	2.74				
PLACE1002737	2.94	0.82	1.29	2.21	2.29	2.41	1.51	1.54	1.54				
PLACE1002741	3.13	1.21	2.06	2.99	2.40	4.38	1.51	3.07	3.07				
PLACE1002752	3.23	1.55	1.75	2.3	2.72	2.29	1.86	1.61	1.61				
PLACE1002763	5.82	2.68	2.79	4.62	5.11	4.63	5.66	4.98	4.98				
PLACE1002766	1.66	0.72	1.60	4.14	2.26	2.27	1.82	1.34	1.34				
PLACE1002772	1.8	1.13	2.05	2.49	1.48	2.20	2.00	2.91	2.91				
PLACE1002782	3.79	1.21	0.99	3.99	3.99	2.22	2.25	2.39	2.39				
PLACE1002784	3.98	1.98	2.41	2.73	2.16	1.89	2.44	4.87	4.87				
PLACE1002798	3.03	1.31	2.50	3.63	5.60	4.46	2.46	3.00	3	*	+		
PLACE1002846	0.71	0.31	1.69	2.44	1.45	2.19	0.63	2.13	2.13				
PLACE1002849	2.59	1.40	2.09	2.06	1.75	1.55	1.88	1.44	1.44				
PLACE1002857	2.54	1.21	2.06	1.63	1.90	1.80	2.01	3.22	3.22				
PLACE1002861	3.24	2.05	2.05	5.01	4.66	4.82	3.10	3.89	3.89	**	+		
PLACE1002872	43.66	21.33	23.44	30.54	23.07	32.80	14.91	18.35	18.35				
PLACE1002877	34.76	13.19	14.79	13.63	20.45	13.77	10.79	13.80	13.8				
PLACE1002879	1.98	0.47	1.85	1.36	3.33	1.12	1.96	1.87	1.87				
PLACE1002886	1.09	0.42	0.92	1.49	1.32	1.87	0.94	1.34	1.34	*	+		
PLACE1002888	3.11	1.53	2.24	1.6	2.71	2.32	1.87	2.30	2.3				
PLACE1002908	4.53	2.06	2.64	3.65	2.87	3.85	3.36	4.12	4.12				
PLACE1002919	5.7	2.20	3.89	5.91	4.05	5.41	4.60	6.30	6.3				
PLACE1002921	1.24	0.74	1.00	0.94	2.00	1.75	1.08	0.94	0.94				
PLACE1002923	2.95	1.00	1.09	2.18	1.25	5.57	0.84	2.57	2.57				
PLACE1002924	4.78	1.22	4.05	2.57	4.25	2.76	1.54	3.00	3				
PLACE1002925	1.27	0.73	0.91	0.45	0.87	0.31	1.52	2.61	2.61	*	+		
PLACE1002931	11.44	4.02	5.58	10.31	11.46	9.16	5.01	7.71	7.71				
PLACE1002935	0.24	0.55	0.45	0.68	0.48	0.50	1.11	1.18	1.18		**	+	
PLACE1002947	4.92	1.59	1.73	2.29	3.03	2.70	3.05	3.68	3.68				
PLACE1002963	1.11	1.73	1.45	1.96	1.63	2.02	1.58	0.96	0.96				
PLACE1002971	2.28	1.16	1.31	3.83	2.51	3.34	2.27	2.89	2.89	*	+		
PLACE1002982	7.21	2.79	4.22	3.07	5.60	7.20	6.47	7.74	7.74				
PLACE1002992	3.36	1.81	0.95	2.29	3.14	1.23	2.29	4.00	4				
PLACE1002995	7.97	4.77	4.17	10.64	14.64	12.97	7.62	12.10	12.1	*	+		
PLACE1002997	3.62	1.37	1.19	4.05	4.29	3.62	2.02	2.74	2.74				
PLACE1010002	3.23	0.90	2.15	1.8	3.62	1.37	1.28	2.45	2.45				
PLACE1010011	3.01	1.89	1.75	1.26	1.49	1.10	1.92	1.85	1.85				
PLACE1010013	1.67	0.86	0.88	1.15	1.55	0.74	1.18	1.56	1.56				
PLACE1010021	2.43	0.87	2.19	2.61	2.46	2.89	3.71	2.58	2.58				
PLACE1010023	4.84	1.80	2.28	2.57	2.67	3.95	1.95	4.34	4.34				
PLACE1010031	5.58	2.99	1.54	4.23	4.90	2.66	3.18	2.93	2.93				
PLACE1010039	1.86	0.50	0.58	0.41	1.70	1.28	1.06	1.35	1.35				

156/180

【0797】

【表326】

1790

PLACE1010045	6.37	3.18	4.06	5.76	9.98	5.99	3.46	7.87	7.87								
PLACE1010053	7.31	4.10	4.89	8.33	10.67	7.68	5.23	4.89	4.89								
PLACE1010060	5.81	2.55	2.85	4.53	3.83	3.76	4.23	4.25	4.25								
PLACE1010069	1.38	1.53	1.33	0.77	1.42	0.61	0.88	2.96	2.96								
PLACE1010070	1.16	0.11	0.64	0.75	0.45	1.16	2.70	1.27	1.27								
PLACE1010074	9.55	3.59	4.51	8.29	7.15	7.46	5.88	9.16	9.16								
PLACE1010076	32.02	14.06	13.18	16.2	20.29	12.88	25.05	26.03	26.03								
PLACE1010078	5.69	2.44	3.34	4.22	3.97	3.39	4.99	6.24	6.24								
PLACE1010081	3.3	1.78	4.36	4.28	4.59	3.29	2.67	5.51	5.51								
PLACE1010083	2.72	1.96	1.66	0.92	1.44	1.20	2.07	2.63	2.63								
PLACE1010089	2.82	1.29	2.28	4.53	3.47	5.64	3.30	4.44	4.44	*		+	*				
PLACE1010096	3.39	1.17	2.00	2.56	2.19	2.70	1.45	1.92	1.92								
PLACE1010102	5.26	3.31	3.97	3.87	6.87	8.63	4.86	8.37	8.37	*		+	*				
PLACE1010105	4.29	0.95	1.09	2.44	2.73	1.94	2.71	4.01	4.01								
PLACE1010106	1.98	0.97	0.87	3.59	2.61	2.19	3.70	3.99	3.99	*		+	**	+			
PLACE1010130	2.14	1.13	1.35	4.01	3.52	4.49	5.26	8.14	8.14	**		+	**	+			
PLACE1010132	6.25	4.26	5.07	4.52	4.25	5.01	4.63	5.39	5.39								
PLACE1010134	3.87	1.25	2.18	2.61	2.68	1.90	2.26	2.82	2.82								
PLACE1010139	28.44	17.86	18.97	14.51	12.81	13.26	30.01	30.01	30.01								
PLACE1010148	2.71	1.27	1.28	1.81	1.73	1.69	1.33	1.07	1.07								
PLACE1010152	2.7	1.53	1.25	4.96	4.00	5.20	3.04	3.45	3.45	*		+	*				
PLACE1010155	1.95	0.77	1.06	1.99	1.84	1.65	3.04	2.97	2.97								
PLACE1010156	1.86	1.01	1.72	5.69	7.58	4.30	7.96	8.94	8.94	*		+	**	+			
PLACE1010161	2.56	0.74	1.26	2.69	3.12	1.69	2.27	2.44	2.44								
PLACE1010181	1.28	0.65	2.02	2	2.26	1.95	1.65	3.46	3.46								
PLACE1010194	4.75	3.52	3.08	5.35	3.77	3.54	4.56	3.30	3.3								
PLACE1010202	1.47	0.70	0.65	1.46	1.47	1.29	1.34	1.69	1.69								
PLACE1010231	1.3	1.19	0.99	2.11	1.60	1.20	1.89	1.43	1.43								
PLACE1010235	2.55	0.79	1.71	2.65	2.39	3.67	1.07	1.49	1.49								
PLACE1010237	0.84	1.17	1.30	1.96	1.56	2.36	1.09	0.99	0.99	*		+					
PLACE1010251	3.81	2.13	2.41	3.72	3.24	1.88	1.45	3.83	3.83								

157/180

【表 3 2 7】

1791

1792

PLACE1010410	4.61	1.87	2.21	5.63	8.60	7.04	3.70	4.91	4.91	*	+		
PLACE1010418	3.29	1.76	2.41	6.21	6.34	6.33	2.79	3.38	3.38	**	+		
PLACE1010425	1.18	0.35	0.46	1.22	0.78	1.70	0.80	1.37	1.37				
PLACE1010443	5.43	3.03	3.71	5.62	3.76	6.48	4.57	5.05	5.05				
PLACE1010445	4.33	2.64	3.67	5.95	5.86	6.97	4.11	3.20	3.2	*	+		
PLACE1010481	1.37	1.21	1.06	0.8	0.77	1.60	1.13	1.10	1.1				
PLACE1010482	5.16	2.61	3.60	3.41	3.22	3.80	5.36	2.91	2.91				
PLACE1010491	2.88	2.21	3.23	5.03	5.64	4.25	5.35	8.41	8.41	*	+	*	+
PLACE1010492	2.47	1.94	1.90	1.59	2.93	3.57	2.66	2.46	2.46				
PLACE1010509	1.31	0.33	0.65	0.44	1.02	0.95	1.07	1.11	1.11				
PLACE1010518	4.3	2.12	3.06	8.55	9.22	8.31	5.08	9.10	9.1	**	+	*	+
PLACE1010522	4.42	3.30	2.99	4.43	3.15	5.70	4.02	5.51	5.51				
PLACE1010529	4.44	3.27	3.34	4.15	2.17	4.43	2.83	4.60	4.6				
PLACE1010547	1.36	0.46	1.84	1.38	2.57	0.83	0.81	0.68	0.68				
PLACE1010560	3.62	1.42	1.78	3.44	4.11	3.17	1.69	3.25	3.25				
PLACE1010562	2.49	1.56	1.51	2.33	1.85	1.73	1.62	1.70	1.7				
PLACE1010579	1.43	1.21	2.19	1.9	1.92	3.18	1.68	1.93	1.93				
PLACE1010580	6.35	2.50	3.66	4.91	4.74	4.81	3.94	5.30	5.3				
PLACE1010599	2.99	2.56	2.79	4.69	2.68	4.02	2.68	2.87	2.87				
PLACE1010606	0.64	1.41	0.70	0.91	1.32	1.04	0.85	0.75	0.75				
PLACE1010616	1.07	0.75	1.12	3.22	1.83	3.57	1.94	1.36	1.36	*	+		
PLACE1010622	9.24	4.26	4.31	2.37	3.79	2.39	1.80	2.04	2.04				
PLACE1010624	6.73	4.32	4.19	2.38	2.68	1.71	1.73	1.83	1.83	*	-	*	-
PLACE1010628	1.26	1.28	1.00	1.32	2.31	1.21	1.10	0.98	0.98				
PLACE1010629	1.86	1.74	1.96	1.86	4.02	4.33	2.68	2.28	2.28		*	+	
PLACE1010630	5.11	3.33	3.71	7.92	7.09	5.39	5.90	7.29	7.29	*	+	*	+
PLACE1010631	1.79	0.95	0.97	2.41	2.47	2.83	1.91	1.86	1.86	*	+		
PLACE1010651	2.68	2.44	2.01	2.53	1.74	2.28	2.68	4.49	4.49				
PLACE1010661	2.42	1.52	2.69	2.28	2.26	4.08	1.65	3.04	3.04				
PLACE1010662	2.49	1.93	2.59	3.46	2.35	2.86	1.94	1.49	1.49				
PLACE1010668	6.55	2.72	2.43	7.07	8.23	6.07	5.21	6.36	6.36				
PLACE1010702	18.26	8.81	10.62	33.41	42.20	27.93	11.82	16.20	16.2	*	+		
PLACE1010709	29.25	14.24	17.35	21.38	21.56	17.73	31.21	41.95	41.95		*	+	
PLACE1010713	11.16	4.98	5.23	9.06	10.03	9.81	9.13	15.19	15.19				
PLACE1010714	0.55	0.48	0.52	0.64	0.75	1.34	0.77	0.70	0.7		**	+	
PLACE1010716	5.99	2.36	2.79	3.78	2.95	5.02	3.07	3.15	3.15				
PLACE1010717	2.06	1.35	1.49	2.22	1.80	2.83	0.90	1.52	1.52				
PLACE1010720	18.67	8.95	8.08	12.05	17.26	10.51	4.13	4.57	4.57				
PLACE1010739	1.36	1.32	0.50	2.03	2.27	3.00	2.05	1.94	1.94	*	+	*	+
PLACE1010743	1.84	1.21	0.69	1.5	0.87	0.37	0.87	1.50	1.5				
PLACE1010752	5.21	2.95	2.72	2.98	3.08	1.69	2.31	3.98	3.98				
PLACE1010761	9.42	7.63	8.64	20.89	19.08	19.20	8.58	11.68	11.68	**	+		
PLACE1010771	7.47	3.15	3.53	5.95	5.91	7.07	6.15	6.64	6.64				
PLACE1010784	0.87	0.52	1.39	0.62	1.10	1.01	1.14	0.89	0.89				
PLACE1010786	3.62	2.60	1.59	2.95	1.86	4.15	2.62	2.64	2.64				
PLACE1010789	2.47	1.71	1.29	7.34	5.69	4.59	3.94	2.83	2.83	**	+		
PLACE1010800	5.09	2.34	2.77	6.42	5.52	4.71	4.26	4.86	4.86				
PLACE1010802	2.85	0.65	1.48	2.19	2.46	1.85	2.00	3.34	3.34				
PLACE1010811	3.15	1.56	1.75	2.32	2.73	2.52	1.52	3.78	3.78				
PLACE1010813	4.37	2.54	2.23	3.08	2.72	2.68	2.51	3.14	3.14				
PLACE1010827	2.09	0.81	0.76	1.38	1.83	1.70	1.14	4.49	4.49				
PLACE1010833	6.2	2.64	3.19	13.01	9.81	9.59	5.31	5.99	5.99	*	+		
PLACE1010839	3.43	2.31	3.38	7.65	5.31	7.09	3.40	4.37	4.37	*	+		
PLACE1010856	3.15	2.01	1.95	2.4	2.08	1.61	2.50	2.16	2.16				
PLACE1010857	5.31	2.37	3.64	2.66	4.62	2.73	2.23	4.31	4.31				
PLACE1010870	6.19	2.76	3.14	7.02	8.56	7.22	4.56	4.25	4.25	*	+		

158/180

【0799】

【表328】

1793

1794

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159/180

【0800】

【表 3 2 9】

1796

[illegible]

160/180

【表 3 3 0】

(900)

特開2002-191363

1797

1798

PLACE1011675	0.49	0.41	0.33	1.54	2.62	1.74	1.03	0.63	0.63	**	+		
PLACE1011682	1.44	1.27	1.77	1.27	1.81	1.50	0.94	2.14	2.14				
PLACE1011708	4.35	4.02	4.14	5.7	7.61	8.08	4.28	4.88	4.88	*	+		
PLACE1011719	1.76	1.55	1.39	2.03	3.35	3.13	2.09	2.66	2.66	*	+	*	+
PLACE1011725	4.47	2.20	1.51	6.52	4.79	5.08	4.70	3.97	3.97				
PLACE1011729	2.26	0.34	1.16	2.9	3.70	2.58	1.88	1.07	1.07	*	+		
PLACE1011741	1.85	1.08	1.46	2.17	2.55	1.44	1.47	2.04	2.04				
PLACE1011749	4.07	1.97	2.35	5.14	5.66	5.55	2.94	2.96	2.96	*	+		
PLACE1011757	7.95	5.78	4.73	28.51	35.97	33.70	18.45	19.91	19.91	**	+	**	+
PLACE1011762	0.6	0.62	0.64	1.51	2.40	1.43	1.31	1.98	1.98	*	+	**	+
PLACE1011778	0.68	0.80	0.85	0.72	1.28	1.91	1.16	0.80	0.8				
PLACE1011783	3.26	3.37	4.33	9.36	8.72	9.68	4.31	3.99	3.99	**	+		
PLACE1011793	2.41	0.78	0.71	3.25	3.16	2.10	1.75	0.51	0.51				
PLACE1011810	1.09	0.35	0.96	0.57	0.96	0.71	1.04	1.41	1.41				
PLACE1011824	1.1	0.61	0.73	1.63	1.19	1.20	1.70	1.61	1.61		**	+	
PLACE1011825	19.56	10.93	11.42	10.37	11.28	11.36	8.08	10.44	10.44				
PLACE1011835	2.12	1.20	1.49	1.76	1.50	1.06	1.49	0.95	0.95				
PLACE1011836	32.53	15.61	18.36	27.63	35.75	28.68	27.23	20.95	20.95				
PLACE1011847	0.74	1.05	0.62	0.62	0.87	1.11	1.60	1.10	1.1				
PLACE1011855	1.16	0.16	0.77	0.69	0.71	1.38	0.70	1.13	1.13				
PLACE1011858	2.38	2.07	1.60	2.19	2.08	1.60	2.79	2.84	2.84		*	+	
PLACE1011874	3.25	1.54	2.03	4.69	4.12	4.23	2.47	3.11	3.11	*	+		
PLACE1011875	1.26	0.66	0.64	1.26	1.14	1.27	0.79	0.74	0.74				
PLACE1011877	6.46	2.58	3.09	3.53	2.30	3.26	2.14	3.12	3.12				
PLACE1011891	1.77	0.88	0.81	1.69	1.67	1.68	1.49	2.31	2.31				
PLACE1011896	0.86	0.25	0.26	0.37	0.26	0.57	0.67	0.48	0.48				
PLACE1011920	2.91	0.83	1.76	1.44	1.22	2.34	1.43	1.54	1.54				
PLACE1011922	4.71	2.40	2.11	4.92	2.79	4.42	3.68	4.23	4.23				
PLACE1011923	3.63	1.24	1.28	5.32	2.65	2.76	7.49	10.90	10.9		**	+	
PLACE1011927	6	2.51	3.82	3.74	4.24	5.24	4.33	4.96	4.96				
PLACE1011939	4.24	2.12	2.87	2.83	3.92	4.33	4.29	5.83	5.83		*	+	
PLACE1011940	5.02	1.82	3.30	7.08	7.36	8.48	4.28	5.85	5.85	*	+		
PLACE1011962	13.26	6.64	7.98	11.22	11.07	13.01	8.70	9.69	9.69				
PLACE1011964	2.09	0.16	0.88	0.97	0.96	0.80	0.82	1.29	1.29				
PLACE1011978	6.83	5.17	5.96	14.23	9.13	15.11	5.12	9.01	9.01	*	+		
PLACE1011980	5.54	2.72	4.54	9.74	12.59	11.01	4.66	6.64	6.64	**	+		
PLACE1011981	6.65	3.37	3.46	5.38	6.10	5.01	3.81	4.69	4.69				
PLACE1011982	0.91	0.32	0.06	0.49	1.81	1.02	0.79	1.02	1.02				
PLACE1011995	4.44	2.50	2.12	5.89	6.15	5.90	3.97	3.81	3.81	*	+		
PLACE1012023	1.79	0.70	1.25	1.43	1.61	2.10	1.24	1.17	1.17				
PLACE1012024	1.87	0.19	0.62	1.01	0.13	0.38	0.66	0.81	0.81				
PLACE1012031	2.22	1.02	2.34	1.31	1.21	3.28	1.23	2.49	2.49				
PLACE2000003	10.16	5.53	7.19	14.74	10.74	18.79	8.25	10.16	10.16				
PLACE2000005	4.58	2.43	2.29	4.4	3.89	4.52	3.11	4.04	4.04				
PLACE2000006	6.31	3.28	0.91	2.52	4.28	3.42	2.99	2.00	2				
PLACE2000007	3.33	1.18	1.86	1.87	3.29	3.13	2.74	4.18	4.18				
PLACE2000011	6.03	2.56	4.49	6.77	5.68	6.77	3.61	4.33	4.33				
PLACE2000014	0.21	0.98	0.82	1.07	2.07	1.53	2.21	2.03	2.03	**	+		
PLACE2000015	1.83	0.65	1.01	1.76	1.29	2.19	1.52	0.85	0.85				
PLACE2000017	3.21	0.73	1.74	6.58	4.80	4.12	2.77	1.87	1.87	*	+		
PLACE2000021	3.22	1.34	2.09	3.94	3.51	4.88	1.82	2.10	2.1	*	+		
PLACE2000022	7.75	3.64	2.82	8.01	9.76	8.90	3.63	6.09	6.09				
PLACE2000030	8.7	4.13	5.99	6.21	7.33	6.10	5.75	6.25	6.25				
PLACE2000032	4.4	0.93	2.84	5.81	6.65	4.83	2.50	4.03	4.03				
PLACE2000033	1.83	1.13	0.57	2.93	3.15	2.60	1.56	1.76	1.76	*	+		
PLACE2000034	2.2	2.03	1.49	1.47	1.71	2.75	1.92	3.95	3.95				

161/180

【0802】

【表331】

1799

1800

PLACE200039	6.48	4.35	4.61	11.87	11.66	13.79	6.80	7.28	7.28	**	+	+
PLACE200040	2.47	1.44	2.20	2.31	3.69	3.32	3.41	4.52	4.52		*	+
PLACE200044	5.02	3.35	3.46	5.51	3.83	5.89	4.93	7.31	7.31			
PLACE200047	8.18	4.36	3.83	9.19	11.31	14.75	5.33	7.74	7.74	*	+	
PLACE200050	12.24	3.78	3.08	8.61	10.29	7.90	7.32	6.64	6.64			
PLACE200061	2.92	0.96	0.97	1.52	0.96	1.26	1.35	1.85	1.85			
PLACE200062	4.77	2.50	2.13	5.58	5.65	5.45	2.96	5.42	5.42	*	+	
PLACE200072	2.7	1.26	2.16	2.17	3.44	2.93	1.74	2.43	2.43			
PLACE200073	1.69	0.72	0.84	1.41	0.59	1.30	1.70	1.52	1.52			
PLACE200077	13.16	8.11	9.49	11.41	12.05	13.08	7.86	8.83	8.83			
PLACE200100	5.14	3.46	2.83	5.96	4.13	5.86	4.27	5.06	5.06			
PLACE200103	4.64	3.10	3.20	7.22	5.44	6.13	4.03	3.95	3.95	*	+	
PLACE200106	7.76	2.85	4.06	6.8	7.28	7.13	4.31	4.99	4.99			
PLACE200111	4.84	2.29	3.47	5	5.26	5.57	4.32	7.27	7.27			
PLACE200115	2.29	0.90	1.18	1.38	0.91	1.85	2.19	2.02	2.02			
PLACE200118	40.98	28.15	29.38	32.74	33.08	38.40	30.44	42.97	42.97			
PLACE200124	16.57	10.11	11.57	19.83	25.65	30.81	16.15	17.74	17.74	*	+	
PLACE200132	7.64	4.32	5.67	5.55	4.79	4.71	7.51	6.44	6.44			
PLACE200136	1.78	0.82	1.05	1.68	1.61	1.41	1.31	1.62	1.62			
PLACE200137	6.66	4.19	3.94	4.2	3.59	5.28	3.96	5.37	5.37			
PLACE200140	9.31	3.10	5.25	7.95	10.19	7.07	4.50	6.74	6.74			
PLACE200147	2.32	1.00	0.75	2.39	2.55	2.14	1.33	2.93	2.93			
PLACE200153	1.79	0.33	0.76	0.89	1.36	1.15	2.17	2.54	2.54	*	+	
PLACE200164	2.92	1.24	1.74	1.97	2.41	1.94	1.21	2.25	2.25			
PLACE200170	4.49	2.57	2.11	5.8	5.33	5.19	3.14	3.80	3.80	*	+	
PLACE200172	3.21	1.40	2.70	1.1	3.14	2.28	1.52	1.72	1.72			
PLACE200173	4.05	3.41	2.95	5.72	7.77	7.43	3.82	4.53	4.53	**	+	
PLACE200174	2.94	1.68	2.28	3.36	3.27	4.06	2.97	2.61	2.61	*	+	
PLACE200176	6.55	2.90	2.44	6.47	6.24	4.58	3.30	4.24	4.24			
PLACE200187	4.34	2.14	1.78	5.63	3.41	5.66	3.80	4.31	4.31			
PLACE200216	4.17	2.38	2.18	6.97	6.14	5.24	7.33	12.03	12.03	*	+	+
PLACE200219	5.75	2.86	2.79	6.33	5.19	5.66	5.15	5.03	5.03			
PLACE200221	6	4.55	4.10	13.08	11.16	10.61	6.14	6.36	6.36	**	+	
PLACE200223	0.66	0.04	0.44	2.56	1.05	0.74	1.35	0.62	0.62			
PLACE200231	2.73	2.97	1.35	3.88	3.66	2.81	3.23	2.76	2.76			
PLACE200235	5.15	3.31	3.10	9.33	15.20	9.28	4.35	5.69	5.69	*	+	
PLACE200246	9.05	5.03	3.92	8.93	10.34	8.27	5.30	6.19	6.19			
PLACE200264	4.4	2.75	1.21	7.23	5.72	5.03	3.18	4.43	4.43	*	+	
PLACE200274	8.27	4.14	5.09	4.88	4.54	3.46	4.83	6.06	6.06			
PLACE200297	14	9.69	10.03	11.98	14.31	14.19	12.42	12.37	12.37			
PLACE200296	3.51	1.96	2.07	2.61	2.73	3.24	2.29	3.69	3.69			
PLACE200302	2.31	2.23	2.10	3.57	4.89	5.77	3.81	3.32	3.32	*	+	+
PLACE200308	7.13	5.46	4.88	12.44	18.75	14.01	6.85	6.47	6.47	*	+	
PLACE200317	1.79	1.81	1.59	2.18	2.88	3.79	2.56	2.49	2.49		**	+
PLACE200324	1.64	0.45	0.66	1.23	0.90	1.15	1.56	1.03	1.03			
PLACE200334	4.7	3.19	3.38	3.36	3.85	3.08	3.51	4.53	4.53			
PLACE200335	6.89	3.67	3.94	9.95	12.98	11.87	4.72	8.12	8.12	**	+	
PLACE200340	1.92	1.00	1.25	2.13	2.37	2.25	1.13	1.70	1.70	*	+	
PLACE200341	4.05	3.76	4.37	3.33	6.79	4.35	3.92	3.97	3.97			
PLACE200342	5.08	6.69	5.71	7.14	7.41	7.08	6.86	8.97	8.97	*	+	+
PLACE200347	4.37	5.20	4.34	7.13	9.07	10.11	6.32	7.54	7.54	*	+	+
PLACE200357	9.87	8.86	7.75	8.78	12.51	9.80	8.70	9.73	9.73			
PLACE200358	4.58	2.20	2.55	4.28	5.06	2.32	5.59	6.09	6.09	*	+	
PLACE200359	2.5	0.52	0.79	3.39	1.81	2.56	2.27	0.82	0.82			
PLACE200366	6.64	3.37	2.29	8.44	9.84	6.44	5.08	3.45	3.45			
PLACE200371	4.65	3.72	1.76	1.73	2.65	2.33	2.69	2.16	2.16			

162/180

【0803】

【表332】

1801

1802

PLACE2000373	4.09	3.75	3.16	3.93	6.78	5.14	3.59	5.16	5.16				
PLACE2000374	3.8	4.38	3.21	5.4	5.00	4.71	4.60	3.34	3.34	*	+		
PLACE2000379	0.43	0.66	0.58	0.91	0.73	1.09	0.79	0.77	0.77	*	+	*	+
PLACE2000386	263.51	193.15	186.41	112.96	134.53	97.90	242.44	237.17	237.2	*	-		
PLACE2000388	6.14	2.57	3.20	4.18	4.37	4.11	3.57	5.67	5.67				
PLACE2000392	22.7	12.68	10.22	19.04	26.24	23.82	20.84	18.58	18.58				
PLACE2000394	4.15	2.33	2.30	7.45	7.62	8.22	3.35	4.27	4.27	**	+		
PLACE2000398	5.77	2.40	4.45	3.51	4.25	5.84	4.07	5.00	5				
PLACE2000399	6.61	3.16	3.15	4.97	4.51	4.35	4.73	5.61	5.61				
PLACE2000402	7.01	4.23	4.20	5.54	4.09	5.56	4.44	3.54	3.54				
PLACE2000404	12.23	7.88	7.30	7.71	7.31	9.74	4.74	6.01	6.01				
PLACE2000411	21.27	11.68	11.82	11.14	10.88	25.73	14.78	18.35	18.35				
PLACE2000418	5.51	3.37	3.01	6.69	5.87	6.09	4.87	3.75	3.75				
PLACE2000419	7.28	4.27	3.30	7.57	9.49	8.40	4.83	4.59	4.59				
PLACE2000425	4.32	2.24	3.29	5.08	4.37	6.06	3.45	3.86	3.86				
PLACE2000427	6.26	3.55	3.23	4.54	4.54	5.08	5.10	5.28	5.28				
PLACE2000433	4.59	2.65	3.36	5.7	5.12	6.87	3.87	4.81	4.81	*	+		
PLACE2000435	29.19	15.24	17.32	14.09	10.07	16.26	23.39	24.72	24.72				
PLACE2000438	3.46	1.48	2.18	3.33	2.20	3.83	3.08	2.95	2.95				
PLACE2000450	9.25	3.49	4.71	9.32	13.42	13.35	5.02	6.24	6.24	*	+		
PLACE2000455	4.87	3.05	1.83	4.35	3.25	3.01	3.72	3.76	3.76				
PLACE2000458	7.14	3.76	3.85	4.27	6.42	5.62	5.42	5.04	5.04				
PLACE2000464	10.07	4.31	6.99	6.94	8.11	6.92	5.43	8.55	8.55				
PLACE2000465	5.73	2.78	3.87	8.13	9.58	9.56	5.26	6.47	6.47	**	+		
PLACE2000473	17.94	8.98	12.76	32.72	23.26	29.31	35.66	50.78	50.78	*	+	**	+
PLACE2000477	1.27	1.02	0.52	1.09	0.78	0.53	1.48	1.22	1.22				
PLACE2000504	7.55	3.19	4.53	8.79	7.24	9.45	5.46	5.75	5.75				
PLACE2000509	61.9	29.47	28.32	32.27	25.30	29.38	45.27	58.28	58.28				
PLACE2000520	9.44	5.05	5.57	6.59	7.39	6.52	4.82	4.55	4.55				
PLACE2000529	9.17	4.67	4.83	9.55	12.07	7.65	6.59	5.44	5.44				
PLACE2000538	3.05	1.65	1.71	3.75	5.45	4.67	2.86	3.09	3.09	*	+		
PLACE2000552	4.37	2.71	2.77	5.23	4.15	6.64	3.13	2.24	2.24				
PLACE2000559	2.05	0.82	1.21	3.28	2.36	2.07	1.89	1.16	1.16				
PLACE2000567	6.3	3.83	5.04	11.45	12.93	15.68	7.26	8.63	8.63	**	+	*	+
PLACE2000569	5.9	3.04	3.53	5	5.11	8.56	5.67	5.68	5.68				
PLACE2000570	27.81	15.78	20.14	32.22	26.02	33.33	21.90	29.50	29.5				
PLACE2000183	2.43	0.95	1.30	3.54	4.94	4.26	1.89	2.90	2.9	*	+		
PLACE2000119	3.74	2.64	1.89	4.89	6.83	4.96	3.78	3.36	3.36	*	+		
PLACE2000121	1.44	1.22	0.45	2.39	2.51	2.11	1.78	2.32	2.32	*	+	*	+
PLACE2000124	5.32	4.70	4.50	12.73	13.74	11.54	6.57	8.87	8.87	**	+	*	+
PLACE2000135	1.71	0.60	0.29	0.53	1.12	0.70	1.32	0.77	0.77	*	+	*	+
PLACE2000136	11.16	7.11	7.56	7.93	9.54	12.38	12.19	8.74	8.74				
PLACE2000142	5.52	1.94	3.53	3.47	2.41	3.28	2.84	4.03	4.03				
PLACE2000149	6.76	3.35	3.37	7.36	7.49	6.80	6.17	8.06	8.06				
PLACE2000147	10.95	4.59	5.26	7.55	8.27	7.90	4.11	4.34	4.34				
PLACE2000148	2.39	0.34	0.58	0.98	1.42	1.44	1.27	2.88	2.88				
PLACE2000154	1.15	0.41	0.41	0.66	1.09	1.15	0.77	2.42	2.42				
PLACE2000155	7.02	4.85	4.71	7.95	8.38	9.70	5.71	8.16	8.16	*	+		
PLACE2000156	15.58	8.96	9.94	8.3	7.38	10.26	12.84	18.30	18.3				
PLACE2000157	4.88	2.44	2.98	2.64	3.32	4.31	4.33	4.39	4.39				
PLACE2000158	8.01	3.79	3.91	9.93	12.06	12.18	6.76	6.67	6.67	*	+		
PLACE2000160	0.65	1.38	1.00	1.99	1.58	2.78	2.19	2.27	2.27			**	+
PLACE2000169	8.16	3.98	2.64	8.16	12.00	9.90	6.56	5.46	5.46				
PLACE2000181	10.19	6.03	5.74	6.99	7.06	6.27	8.77	10.50	10.5				
PLACE2000194	3.17	1.81	1.47	3.38	2.68	2.61	2.86	3.96	3.96				
PLACE2000197	0.71	0.07	1.20	1.17	0.86	1.14	4.41	2.01	2.01				

163/180

【0804】

【表333】

1804

PLACE3000199	3.29	1.08	1.38	2.04	1.59	1.81	1.36	3.52	3.52				
PLACE3000205	9.93	4.59	5.70	17.83	17.57	18.45	14.66	13.74	13.74	**	+	*	+
PLACE3000207	5.7	3.47	2.72	7.85	6.73	9.27	4.82	3.93	3.93	*	+		
PLACE3000208	5.91	3.83	2.56	4.66	4.50	5.84	3.33	5.31	5.31				
PLACE3000213	3.26	1.41	0.88	1.83	1.88	1.34	1.39	1.20	1.2				
PLACE3000215	5.27	3.26	2.05	2.91	1.77	2.17	4.16	5.65	5.65				
PLACE3000218	0.67	1.20	0.52	0.53	0.72	1.11	0.94	1.60	1.6				
PLACE3000220	4.81	2.27	2.38	5.89	5.17	5.82	4.14	4.16	4.16	*	+		
PLACE3000221	18.58	12.33	11.49	19.49	17.73	21.75	11.62	11.46	11.46				
PLACE3000225	2.26	1.52	1.43	2.24	4.06	3.45	1.47	2.45	2.45				
PLACE3000226	4.27	2.49	2.02	2.27	5.71	4.75	1.91	2.73	2.73				
PLACE3000230	2.53	2.38	1.81	1.66	1.64	1.71	2.48	1.35	1.35				
PLACE3000231	3.29	1.13	0.60	2.47	2.81	2.21	3.05	2.05	2.05				
PLACE3000235	3.68	1.67	2.02	7.18	5.86	5.62	2.96	4.70	4.7	**	+		
PLACE3000242	4.95	3.58	3.28	11.36	12.88	9.51	10.16	9.35	9.35	**	+	**	+
PLACE3000244	1.78	1.29	0.91	1.71	1.41	0.91	1.35	0.85	0.85				
PLACE3000253	1.86	1.24	1.41	3.62	2.97	3.37	3.19	2.28	2.28	**	+	*	+
PLACE3000254	51.54	34.63	40.51	40.03	46.12	56.93	50.43	47.16	47.16				
PLACE3000271	5.35	3.90	4.49	12.52	15.43	16.28	5.75	8.41	8.41	**	+	*	+
PLACE3000276	1.34	1.63	0.94	1.51	1.84	1.69	1.54	1.70	1.7				
PLACE3000304	29.17	18.78	18.07	35.22	34.12	39.27	19.90	28.29	28.29	*	+		
PLACE3000309	5.85	2.02	1.54	4.32	5.65	5.33	3.03	4.10	4.1				
PLACE3000310	2.86	0.49	0.75	1.95	1.51	1.29	0.96	1.26	1.26				
PLACE3000320	2.43	0.72	1.39	2.35	2.57	2.63	2.57	2.39	2.39				
PLACE3000332	3.17	2.14	2.01	4.49	4.42	5.13	3.26	3.42	3.42	**	+		
PLACE3000330	3.98	4.24	5.26	4.75	5.64	8.32	9.28	8.32	8.32			**	+
PLACE3000331	3.82	3.74	4.92	7.37	8.26	9.20	3.96	4.94	4.94	**	+		
PLACE3000334	2.26	2.25	2.90	3.09	4.08	3.48	1.74	3.42	3.42	*	+		
PLACE3000339	1.51	1.25	0.97	2.83	3.03	1.44	3.34	1.37	1.37				
PLACE3000341	4.76	1.28	0.07	6.03	6.07	5.79	3.01	2.61	2.61	*	+		
PLACE3000350	3.67	2.80	1.30	3.28	4.94	3.47	3.39	2.10	2.1				
PLACE3000352	6.08	5.05	2.30	5.5	5.48	4.50	3.98	4.70	4.7				
PLACE3000353	0.84	1.44	1.91	1.76	2.65	2.70	3.03	3.61	3.61		**	+	
PLACE3000362	1.98	1.66	1.84	6.16	5.62	6.95	2.53	2.39	2.39	**	+	**	+
PLACE3000363	0.72	2.27	1.87	2.22	2.71	1.75	1.29	1.32	1.32				

164/180

1805

1806

FLACE000014	5.92	2.92	3.44	5.18	6.07	5.84	4.46	4.89	4.89							
FLACE000039	1.91	1.44	1.35	3.21	1.93	3.26	3.99	3.79	3.79					**	+	
FLACE000034	2.6	1.30	1.44	3.92	3.82	4.60	4.01	3.41	3.41	**	+	*	*	*		
FLACE000049	10.4	5.48	5.72	12.83	16.95	11.80	9.94	9.10	9.1	*	*		*	*		
FLACE000052	6.49	3.73	2.47	4.77	4.77	5.30	5.23	5.62	5.62							
FLACE000062	6.59	2.48	4.03	4.7	5.26	5.48	4.59	4.62	4.62							
FLACE000063	7.7	3.50	3.52	6.91	6.71	9.08	5.77	5.40	5.4							
FLACE000089	2.96	1.45	2.33	5.97	4.11	5.63	4.54	4.57	4.57	*	+	**	*			
FLACE000093	2.81	1.09	0.89	1.95	1.62	1.17	2.18	1.71	1.71							
FLACE000100	4.42	2.89	2.49	3.93	3.32	5.21	3.23	2.62	2.62							
FLACE000103	5.02	1.97	1.98	3.66	2.71	3.95	2.81	2.33	2.33							
FLACE000106	8.72	4.11	3.74	4.38	5.75	4.55	4.28	4.16	4.16							
FLACE000128	7.39	4.68	3.31	9.85	9.72	8.43	7.44	6.38	6.38	*	+					
FLACE000129	6.04	2.07	2.84	4.76	6.70	6.24	4.40	2.79	2.79							
FLACE000131	8.08	5.12	4.57	12.93	9.62	6.75	8.38	9.08	9.08							
FLACE000147	1.54	0.95	0.56	0.28	1.32	1.44	1.32	1.12	1.12							
FLACE000156	10.36	6.90	8.62	23.53	13.89	24.29	10.99	14.64	14.64	*	+					
FLACE000175	2.77	1.36	1.67	3	2.23	3.75	2.99	2.63	2.63							
FLACE000190	25.73	14.17	16.07	19.71	16.55	18.77	20.04	22.67	22.67							
FLACE000192	19.18	10.39	8.86	17.39	19.36	14.48	12.50	10.81	10.81							
FLACE000206	26.35	11.24	12.17	18.68	19.88	13.96	10.44	9.28	9.28							
FLACE000211	17.59	9.35	9.22	14.45	14.14	14.09	11.01	11.86	11.86							
FLACE000214	3.16	2.15	2.41	4.6	3.22	2.93	3.58	2.23	2.23							
FLACE000223	5.13	3.77	3.41	7.67	6.23	6.64	5.04	5.14	5.14	*	+					
FLACE000223	5.15	2.40	3.83	4.77	3.40	3.75	4.17	5.28	5.28							
FLACE000229	2.61	1.29	1.59	3.13	1.82	2.66	3.16	3.28	3.28				*	+		
FLACE000230	10.54	4.47	5.13	3.92	4.50	6.23	2.12	1.74	1.74							
FLACE000233	7.43	4.11	1.84	9.98	7.86	6.99	4.69	5.82	5.82							
FLACE000239	10.37	3.20	3.64	8.75	7.61	7.98	4.24	5.32	5.32							
FLACE000247	3.98	2.15	1.70	4.78	4.11	3.53	4.31	3.20	3.2							
FLACE000250	6.06	3.58	4.71	8.33	8.43	6.31	5.56	7.08	7.08	*	+					
FLACE000251	2.91	1.12	1.52	2.79	1.94	3.45	2.33	2.20	2.2							
FLACE000259	8.04	3.19	7.29	6.61	5.24	7.03	5.35	5.02	5.02							
FLACE000261	12.86	7.43	11.27	7.94	6.30	11.29	13.49	12.71	12.71							
FLACE000264	5.07	2.86	1.88	6.35	6.52	5.02	3.87	4.16	4.16							
FLACE000269	8.57	4.36	5.52	8.01	9.34	7.35	6.32	5.77	5.77							
FLACE000270	3.13	1.82	0.87	2.42	1.82	3.08	1.61	2.16	2.16							
FLACE000281	19.68	7.73	9.21	20.75	31.26	26.50	19.08	19.52	19.52	*	+					
FLACE000380	6.08	3.69	5.60	7.08	6.91	5.29	4.32	5.19	5.19							
FLACE000320	5.62	3.77	3.47	7.13	6.02	6.80	4.81	4.30	4.3	*	+					
FLACE000323	8.19	5.61	3.78	9.71	7.40	10.97	6.79	7.01	7.01							
FLACE000326	4.48	1.87	1.75	4.11	3.23	4.42	3.33	2.91	2.91							
FLACE000344	2.79	2.15	2.50	2.98	1.69	2.74	1.96	2.31	2.31							
FLACE000347	20.7	10.82	8.58	19.27	12.61	11.57	8.40	11.08	11.08							
FLACE000354	4.74	1.02	1.75	4.04	3.76	1.42	1.52	3.10	3.1							
FLACE000367	2.52	1.53	1.38	2.65	2.91	2.67	2.13	2.39	2.39							
FLACE000369	4.83	2.57	3.31	5.06	4.32	4.54	4.37	5.39	5.39							
FLACE000379	5.69	3.46	3.43	6.11	6.24	7.35	4.30	5.61	5.61	*	+					
FLACE000387	3.69	1.95	1.27	2.17	3.32	3.40	2.28	3.13	3.13							
FLACE000392	1.14	0.56	0.17	0.91	1.44	1.78	1.07	1.00	1							
FLACE000399	23.89	17.50	15.29	24.02	23.23	28.14	19.33	22.84	22.84							
FLACE000401	1.48	0.84	0.45	2.03	4.11	1.07	1.24	1.78	1.78							
FLACE000403	9.89	5.20	5.81	9.29	8.13	6.25	5.57	7.91	7.91							
FLACE000411	5.72	2.12	2.75	5.81	4.30	5.15	4.86	3.29	3.29							
FLACE000415	3.21	2.22	2.69	3.67	4.43	3.44	4.57	6.28	6.28				**	+		
FLACE000416	4.63	3.13	2.08	5.57	4.05	4.56	5.50	4.53	4.53							

165/180

【0806】

【表 3 3 5】

1809

1810

SKNMC1000099	4.27	1.82	4.32	2.68	2.85	4.25	4.90	2.31	2.31										
SKNMC1000104	2.88	1.34	1.64	2.26	2.75	3.25	1.82	2.06	2.06										
SKNMC1000113	2.91	1.98	1.70	2.53	3.12	2.50	2.17	2.08	2.08										
SKNMC1000119	4.61	2.84	2.09	3.6	4.44	4.19	3.90	3.35	3.35										
SKNMC1000143	2.86	0.96	0.73	2.73	1.96	2.31	2.39	2.51	2.51										
SKNMC1000170	4.02	1.58	1.54	3.23	3.13	3.73	2.53	3.66	3.66										
SKNMC1000178	5.92	3.14	3.92	5.65	4.47	6.23	4.68	4.57	4.57										
SKNMC1000194	3.57	2.37	1.14	2.02	1.84	1.46	1.82	1.68	1.68										
SKNMC1000198	4.86	3.19	3.66	3.95	2.35	5.30	3.90	3.61	3.61										
SKNMC1000225	3.86	1.48	1.25	3.04	2.83	3.41	1.69	1.50	1.5										
SKNMC1000249	2.6	1.16	0.14	2.11	0.98	1.05	0.97	1.03	1.03										
SPLN1000007	3.1	1.45	1.01	2.61	2.77	3.19	1.50	2.71	2.71										
SPLN1000012	4.58	1.70	1.35	3.53	2.59	2.41	3.41	4.25	4.25										
SPLN1000014	6.11	2.53	3.00	5.55	7.51	4.48	3.02	3.02	3.02										
SPLN1000036	2.67	1.59	1.60	2.81	3.21	2.90	3.30	2.69	2.69										
SPLN1000059	0.04	0.28	0.35	0.37	0.20	0.93	0.51	0.65	0.65								*	+	
SPLN1000068	2.47	1.01	1.48	3.14	3.20	4.62	4.16	2.46	2.46								+		
SPLN1000072	3.94	2.95	2.34	4.26	4.36	3.28	3.61	3.41	3.41										
SPLN1000101	41.57	16.82	24.85	23.9	21.81	9.24	15.06	12.84	12.84										
SPLN1000108	3.06	1.50	1.01	2.01	2.01	1.31	1.57	2.16	2.16										
SPLN1000113	4.35	2.46	2.67	4.83	2.55	2.28	3.11	3.66	3.66										
SPLN1000114	2.42	2.37	1.43	3.43	2.78	2.56	2.74	3.97	3.97								*	+	
SPLN1000132	4.91	2.27	3.07	3.65	2.33	4.08	4.07	4.65	4.65										
SPLN1000135	4.83	1.59	3.15	4.45	2.38	2.83	5.59	5.94	5.94								*	+	
SPLN1000136	4.48	3.01	2.79	7.59	5.71	8.15	9.03	12.90	12.9								+	**	+
SPLN1000141	2.18	1.15	1.72	2.22	2.60	2.27	2.35	1.59	1.59										
SPLN1000164	4.46	1.47	1.76	5.13	4.33	4.86	3.29	5.58	5.58										
SPLN1000166	2.49	0.67	1.05	2.36	3.89	2.42	2.08	3.68	3.68										
SPLN1000175	5.45	3.05	4.54	4.81	4.46	4.23	3.32	5.47	5.47										
SPLN1000182	2.6	0.65	0.61	1.52	1.41	2.22	1.31	1.69	1.69										
SPLN1000185	3.66	1.87	1.77	5.3	4.71	4.35	5.29	7.02	7.02								*	**	+
THYMU1000004	14.86	7.77	9.02	24.57	18.18	21.23	10.89	18.76	18.76								+		
THYMU1000009	8.45	5.32	5.87	7.04	5.33	4.60	6.33	5.23	5.23										
THYMU1000015	26.6	19.78	21.97	16.29	13.38	16.01	9.72	8.42	8.42								*	**	-
THYMU1000016	8.26	4.04	3.89	15.26	18.83	11.55	9.39	7.02	7.02								+		
THYMU1000023	3.89	1.34	1.23	2.77	2.08	3.06	2.39	2.39	2.39										
THYMU1000034	2.61	1.47	0.66	2.74	1.63	1.39	1.31	3.64	3.64										
THYMU1000035	1.07	0.61	0.61	0.44	0.64	0.76	1.85	2.01	2.01								**	+	
THYMU1000037	1.82	1.82	1.19	2.22	2.35	0.98	2.22	2.11	2.11										
THYMU1000042	10.49	6.31	8.55	6.35	4.98	6.18	8.88	5.36	5.36										
THYMU1000047	4.11	2.46	3.11	10.3	9.57	11.11	4.37	4.74	4.74								**	+	+
THYMU1000080	3.32	3.11	1.09	3.11	4.52	4.74	2.28	1.83	1.83										
THYMU1000094	32.63	25.01	18.12	54.59	42.21	15.66	23.80	19.03	19.03										
THYMU1000109	8.44	4.34	3.79	6.74	8.15	5.93	7.23	6.42	6.42										
THYMU1000127	6.78	3.40	3.18	8.92	8.62	7.88	6.21	6.83	6.83								*	+	
THYMU1000130	4.13	1.20	1.02	4.32	4.36	3.32	2.41	3.21	3.21										
THYMU1000137	4.62	2.65	2.71	3.35	5.77	3.60	4.29	4.56	4.56										
THYMU1000146	4.71	3.58	4.49	7.3	4.71	6.30	5.63	4.56	4.56										
THYMU1000159	25.83	22.19	24.81	10.37	8.83	13.56	14.02	11.77	11.77								**	**	-
THYMU1000163	6.99	6.16	7.74	9.39	10.76	10.43	5.92	8.13	8.13								**	+	
THYMU1000167	2.34	1.29	1.70	2.93	3.51	2.43	1.61	1.52	1.52										
THYMU1000186	5.07	2.17	2.10	3.12	3.14	2.53	2.77	2.94	2.94										
THYRO1000017	5.52	2.12	2.08	4.91	6.19	6.15	4.38	3.34	3.34										
THYRO1000026	3.58	2.32	1.61	2.83	7.78	2.86	6.59	2.67	2.67										
THYRO1000034	3.17	3.08	1.76	3.93	4.73	4.25	3.60	4.07	4.07								*	+	
THYRO1000035	1.48	0.66	0.72	1.53	1.40	2.82	0.67	2.32	2.32										

167/180

【0808】

【表337】

1811

1812

THYRO1000036	1.47	2.88	1.52	4.59	3.60	4.55	3.37	2.09	2.09	*	+		
THYRO1000040	3.94	3.60	4.34	8.08	4.14	6.42	4.83	5.15	5.15		*		+
THYRO1000061	5.94	3.66	2.97	4.84	6.31	3.19	4.68	3.52	3.52				
THYRO1000067	15.2	9.77	9.78	10.78	13.27	13.10	11.70	12.59	12.59				
THYRO1000070	6	3.76	5.68	6.21	9.32	7.75	5.41	6.34	6.34				
THYRO1000072	2.94	1.82	1.84	5.83	8.39	3.32	2.14	2.54	2.54				
THYRO1000084	4.5	1.85	2.58	3.76	4.67	3.19	3.46	2.16	2.16				
THYRO1000085	10.88	11.54	13.23	14.79	17.02	16.91	12.99	15.14	15.14	*	+		
THYRO1000086	0.12	1.27	1.00	1.39	0.92	1.37	0.61	1.10	1.1				
THYRO1000087	0.56	0.67	0.91	1.37	1.09	0.98	1.47	0.51	0.51	*	+		
THYRO1000092	6	2.56	1.98	8.27	6.56	7.42	3.48	3.45	3.45	*	+		
THYRO1000093	1.44	1.12	0.93	2.32	0.88	2.13	1.21	1.43	1.43				
THYRO1000099	5.17	1.21	1.50	4.31	3.36	5.55	2.12	3.30	3.3				
THYRO1000107	2.2	0.53	1.13	2.82	7.80	4.79	2.15	2.47	2.47				
THYRO1000111	1.83	0.33	0.78	2.31	3.19	3.86	1.66	1.58	1.58	*	+		
THYRO1000121	3.44	1.10	1.03	3.02	3.40	6.52	2.38	1.76	1.76				
THYRO1000124	2.37	0.51	0.78	3.06	2.51	2.25	0.89	1.60	1.6				
THYRO1000129	1.3	0.82	0.51	1.26	1.52	1.53	0.49	1.02	1.02				
THYRO1000130	3.62	2.11	2.49	5.24	10.43	5.75	7.64	2.92	2.92				
THYRO1000132	8.41	1.76	1.74	4.45	6.81	7.03	2.87	3.11	3.11				
THYRO1000134	3.55	1.81	2.95	6.64	4.07	4.40	3.58	4.01	4.01				
THYRO1000144	13.82	5.38	3.94	8.01	7.60	7.93	4.07	4.00	4				
THYRO1000155	2.5	0.51	0.58	3.49	1.11	0.97	0.53	1.08	1.08				
THYRO1000156	1.89	1.44	0.82	2.61	2.67	3.19	1.97	1.97	1.97	*	+		
THYRO1000163	3.98	1.47	3.15	9.1	7.23	11.51	7.86	4.19	4.19	*	+		
THYRO1000173	2.9	2.72	1.68	4.44	4.27	4.08	1.67	3.61	3.61	**	+		
THYRO1000186	9.1	5.19	4.20	10.2	15.51	9.61	7.74	7.44	7.44				
THYRO1000187	5.63	2.01	3.20	6.21	7.01	6.32	5.05	3.18	3.18				
THYRO1000190	2.89	1.46	2.17	5.4	4.76	5.31	4.40	2.66	2.66	**	+		
THYRO1000196	0.92	0.80	1.33	2.19	1.72	1.35	0.94	1.18	1.18				
THYRO1000197	3.18	2.33	2.51	5.88	3.71	6.16	4.77	4.51	4.51	*	+	**	+
THYRO1000199	3.03	1.48	1.85	2.3	1.87	3.05	2.39	2.56	2.56				
THYRO1000206	14.52	5.55	4.65	11.65	9.64	12.12	6.54	6.11	6.11				
THYRO1000211	5.01	1.90	2.05	5.6	6.77	7.34	2.67	3.86	3.86	*	+		
THYRO1000213	7.73	2.24	1.94	3.18	4.68	4.24	4.78	2.83	2.83				
THYRO1000218	1.72	0.91	0.91	5.64	4.49	4.50	3.42	4.40	4.4	**	+	**	+
THYRO1000241	3.26	1.56	2.99	5.29	5.78	7.35	4.01	4.55	4.55	*	+	*	+
THYRO1000243	6.01	2.48	2.81	8.74	10.47	5.58	3.38	6.54	6.54				
THYRO1000246	2.49	0.94	1.13	2.44	2.95	2.72	4.13	4.49	4.49			**	+
THYRO1000253	3.03	2.39	2.12	4	3.56	6.64	2.33	3.27	3.27				
THYRO1000270	0.85	0.93	0.64	2.95	1.36	0.98	0.55	0.45	0.45		*		
THYRO1000272	2.19	0.22	0.27	0.43	1.67	1.01	0.46	0.46	0.46				
THYRO1000285	6.19	3.69	1.88	4.45	4.41	6.07	8.92	4.83	4.83				
THYRO1000288	7.58	2.67	2.64	4.38	5.78	3.82	4.63	6.75	6.75				
THYRO1000296	3.95	2.07	1.83	3.07	4.49	3.23	3.68	3.54	3.54				
THYRO1000320	4.13	0.95	0.96	3.75	3.95	6.99	3.33	5.20	5.2				
THYRO1000322	38.05	21.86	30.50	21.36	20.13	23.75	18.89	19.42	19.42				
THYRO1000327	1.02	0.47	0.74	3.44	1.87	3.51	2.40	2.03	2.03	*	+	**	+
THYRO1000343	3.18	0.96	1.50	2.27	1.34	2.27	1.96	1.19	1.19				
THYRO1000345	4.6	2.12	2.05	3.98	5.40	3.79	1.33	1.91	1.91				
THYRO1000358	7.71	5.28	3.61	7.26	4.42	4.45	5.44	7.71	7.71				
THYRO1000360	11.25	3.81	3.69	7.91	6.70	6.11	5.37	4.82	4.82				
THYRO1000375	6.52	5.33	3.32	11.74	11.72	9.07	7.25	13.34	13.34	*	+	*	+
THYRO1000381	1.08	0.73	0.85	2.03	1.76	1.73	1.91	1.07	1.07	**	+		
THYRO1000387	2.85	2.46	2.45	4.71	4.58	5.18	3.81	2.92	2.92	**	+		
THYRO1000394	3.11	2.36	2.61	4.86	4.51	5.33	6.21	6.15	6.15	**	+	**	+

168/180

【0809】

【表338】

1813

1814

THYRO1000395	4.25	2.93	1.91	4.03	3.11	3.93	4.17	2.18	2.18										
THYRO1000400	4.41	1.20	1.12	2.44	2.11	3.30	1.51	2.67	2.67										
THYRO1000401	5.78	2.72	2.22	4.86	5.69	4.69	3.46	3.98	3.98										
THYRO1000407	2.85	1.30	0.87	2.33	1.72	1.87	2.55	3.06	3.06										
THYRO1000420	6.84	3.72	3.92	6.3	4.99	6.57	4.27	4.92	4.92										
THYRO1000438	3.47	2.61	5.10	3.55	4.73	5.14	3.74	2.32	2.32										
THYRO1000453	3.79	2.27	3.32	4.32	3.39	3.80	3.50	2.68	2.68										
THYRO1000455	0.86	0.19	0.08	0.98	0.97	1.02	0.43	0.69	0.69										
THYRO1000471	3.13	0.99	1.71	4.82	2.11	3.45	2.03	2.21	2.21										
THYRO1000481	3.05	2.09	1.78	2.49	2.59	3.24	2.75	3.65	3.65										
THYRO1000484	7.3	2.87	2.29	10.67	15.51	6.38	4.46	3.81	3.81										
THYRO1000488	1.1	0.92	1.15	1.45	1.81	1.35	2.24	2.38	2.38	*	+	**	+						
THYRO1000501	2.42	1.63	1.50	2.59	2.38	2.19	2.65	3.01	3.01										
THYRO1000502	1.72	1.26	1.14	1.06	1.74	2.09	1.25	1.88	1.88										
THYRO1000505	1.86	1.15	0.80	1	1.66	1.13	1.93	1.49	1.49										
THYRO1000535	3.34	1.94	2.04	4.99	3.71	3.63	10.07	9.11	9.11										
THYRO1000556	3.48	3.02	2.08	3.02	2.21	3.79	3.38	3.27	3.27										
THYRO1000558	2.31	1.23	1.10	1.93	1.95	2.49	2.30	1.39	1.39										
THYRO1000569	37.42	23.06	26.88	28.52	31.17	30.05	27.41	43.25	43.25										
THYRO1000579	3.86	2.04	1.70	2.58	2.40	4.33	2.86	3.78	3.78										
THYRO1000572	2.15	0.94	1.24	2.2	1.78	1.73	2.48	3.26	3.26										
THYRO1000573	2.15	0.40	1.11	1.23	2.42	1.75	1.79	2.04	2.04										
THYRO1000577	1.28	1.14	0.64	1.15	1.13	1.55	1.85	1.41	1.41										
THYRO1000580	5.42	3.17	3.10	6.46	6.34	9.14	4.00	4.26	4.26	*	+								
THYRO1000584	2.72	2.07	1.38	2.78	3.98	3.94	2.67	3.22	3.22										
THYRO1000585	2.25	1.51	1.61	5.52	5.02	4.69	3.92	4.40	4.40	**	+	**	+						
THYRO1000596	0.84	0.25	0.33	0.85	1.98	1.44	1.19	1.17	1.17										
THYRO1000602	5.45	3.58	2.07	8.38	7.15	5.61	4.80	5.98	5.98										
THYRO1000605	3.06	1.73	1.76	2.38	1.83	1.39	2.18	2.05	2.05										
THYRO1000615	1.88	0.80	0.63	1.19	1.72	1.17	1.04	2.25	2.25										
THYRO1000625	3.03	2.54	1.58	4.59	3.95	5.93	3.48	4.60	4.60	*	+	*	+						
THYRO1000636	2.66	2.57	2.75	6.51	3.94	8.33	4.69	4.10	4.10	*	+	**	+						
THYRO1000637	1.23	0.82	0.65	1.88	1.42	1.92	2.10	1.39	1.39	*	+								
THYRO1000641	1.4	0.60	1.08	0.89	1.31	1.56	1.11	0.84	0.84										
THYRO1000657	3.65	3.07	3.41	3.91	3.79	3.12	1.96	2.62	2.62										
THYRO1000658	7.81	3.42	3.03	11.25	11.55	11.93	5.08	5.90	5.90	*	+								
THYRO1000662	2.88	1.16	0.83	2.17	1.76	1.90	1.97	1.81	1.81										
THYRO1000666	2.42	0.88	1.16	3.25	2.79	4.33	1.98	2.43	2.43	*	+								
THYRO1000676	2.32	1.10	0.52	2.88	3.21	3.68	3.68	2.15	2.15	*	+								
THYRO1000678	-0.09	0.33	0.95	0.54	0.74	1.28	1.19	2.92	2.92										
THYRO1000684	1.03	2.45	1.63	3.34	3.15	3.52	4.80	2.39	2.39	*	+								
THYRO1000694	2.71	3.51	4.23	5.53	5.34	4.52	4.35	3.80	3.80	*	+								
THYRO1000699	15.82	15.18	11.44	15.15	15.90	16.09	15.44	10.86	10.86										
THYRO1000712	3.39	2.96	2.14	8.58	5.42	7.84	3.11	4.20	4.20	*	+								
THYRO1000715	4.02	2.34	2.31	2.86	4.26	2.85	3.39	2.68	2.68										
THYRO1000716	2.32	0.65	1.04	2.97	4.44	2.89	2.03	1.56	1.56	*	+								
THYRO1000717	2.15	0.84	1.30	4.23	5.94	4.84	1.47	3.93	3.93	**	+								
THYRO1000723	0.84	0.47	0.25	0.76	1.49	1.41	0.88	0.44	0.44										
THYRO1000734	0.78	0.54	0.33	1.36	1.43	0.84	0.50	0.83	0.83	*	+								
THYRO1000748	0.59	2.46	1.89	4.51	7.18	3.35	2.76	2.25	2.25										
THYRO1000755	6.84	4.25	3.30	14.94	19.03	9.44	6.39	7.81	7.81	*	+								
THYRO1000756	3.41	1.39	1.44	2.12	2.71	3.18	1.98	2.77	2.77										
THYRO1000776	1.32	1.08	1.00	2.41	2.21	1.88	2.52	1.74	1.74	**	+	*	+						
THYRO1000777	2.84	1.43	1.39	4.03	2.36	3.89	2.12	2.28	2.28										
THYRO1000779	0.67	0.56	0.16	1.05	0.79	0.60	0.44	0.25	0.25										
THYRO1000782	3.17	1.32	2.40	4.64	3.68	4.39	4.70	5.63	5.63	*	+	**	+						

169/180

【0810】

【表339】

1815

1816

THYRO1000783	1.63	0.89	1.30	3.1	2.26	1.53	1.80	1.30	1.3										
THYRO1000786	4.89	2.61	2.30	6.28	3.05	5.87	4.15	4.10	4.1										
THYRO1000787	10.6	5.80	4.42	7.07	6.40	5.00	7.52	6.30	6.3										
THYRO1000792	6.58	1.87	1.67	2.34	3.23	1.91	2.22	2.34	2.34										
THYRO1000793	2.04	0.81	0.90	2.24	3.46	2.95	1.63	1.90	1.9*	+									
THYRO1000795	2.76	1.16	1.46	2.99	2.52	3.49	2.58	3.17	3.17										
THYRO1000796	2.38	0.64	1.44	4.8	3.84	4.16	2.52	2.59	2.59**	+									
THYRO1000798	3.16	1.83	2.57	4.6	3.74	3.94	2.76	3.06	3.06*	+									
THYRO1000800	7.44	4.89	4.90	15.03	11.25	16.69	6.56	6.96	6.96**	+									
THYRO1000805	0.7	1.04	0.84	1.39	1.41	1.19	1.16	1.27	1.27*	+									
THYRO1000815	7	4.02	3.01	10.69	12.71	10.92	7.46	5.49	5.49**	+									
THYRO1000829	4.85	1.50	0.99	3.49	4.27	2.08	2.62	2.36	2.36										
THYRO1000835	2.11	1.21	1.15	2.86	3.23	3.63	2.50	4.32	4.32*	+									
THYRO1000843	5.05	2.38	2.97	4.77	5.02	6.46	4.36	3.37	3.37										
THYRO1000846	2.51	1.06	0.98	2.34	1.74	1.56	2.17	1.43	1.43										
THYRO1000853	2.42	0.77	2.13	2.09	1.40	2.69	3.08	3.10	3.1										
THYRO1000855	4.5	4.43	3.85	5.88	4.56	7.12	5.76	3.18	3.18										
THYRO1000865	3.16	2.10	3.34	4.86	6.09	6.43	5.14	2.65	2.65**	+									
THYRO1000866	11.62	9.40	6.30	9.67	9.65	5.08	11.39	9.54	9.54										
THYRO1000881	36.03	18.32	15.54	24.61	23.19	29.23	22.14	28.98	28.98										
THYRO1000894	3.99	1.72	1.92	2.01	2.07	2.23	2.83	2.03	2.03										
THYRO1000895	2.03	0.86	1.43	1.55	2.22	2.83	1.11	1.40	1.4										
THYRO1000916	3.35	1.86	1.68	6.43	4.60	5.32	3.15	2.84	2.84*	+									
THYRO1000917	19.78	13.58	15.27	18.14	13.63	19.21	15.55	24.10	24.1										
THYRO1000926	3.79	1.84	2.71	4.53	2.38	2.98	3.39	2.18	2.18										
THYRO1000934	0.9	1.09	0.59	2.64	2.45	2.04	2.64	2.12	2.12**	+									
THYRO1000951	4.53	2.89	1.88	3.09	4.97	2.59	3.21	3.92	3.92										
THYRO1000952	3.27	1.18	1.32	2.44	2.17	2.23	1.41	2.31	2.31										
THYRO1000954	2.11	1.50	1.47	2.05	2.05	1.60	2.11	2.25	2.25										
THYRO1000960	5.02	0.63	1.57	3.83	4.64	3.41	3.77	4.16	4.16										
THYRO1000961	1.21	1.05	0.73	2.4	1.40	1.52	2.97	2.62	2.62										
THYRO1000964	2.36	2.00	1.45	3.05	2.41	3.11	3.20	2.63	2.63										
THYRO1000971	6.39	3.74	2.87	7.64	6.60	7.93	4.97	5.58	5.58										
THYRO1000974	8.5	6.07	6.15	9.83	9.20	11.43	9.21	8.90	8.9*	+									
THYRO1000975	6.08	2.45	2.54	7.25	6.73	7.67	5.66	3.65	3.65*	+									
THYRO1000983	6.75	2.78	2.84	5.03	3.45	3.63	5.16	7.50	7.5										
THYRO1000984	4.73	2.02	2.56	6.84	6.78	4.19	3.85	4.94	4.94										
THYRO1000988	5.73	4.61	2.66	9.09	5.83	6.82	5.38	4.73	4.73										
THYRO1000991	5.53	2.99	3.68	7.73	4.24	7.52	5.28	4.92	4.92										
THYRO1000999	1.49	2.18	1.52	3.22	3.15	4.39	2.64	2.87	2.87*	+									
THYRO1001003	3.32	1.87	1.67	2.91	2.45	1.95	2.38	1.98	1.98										
THYRO1001015	6.07	3.22	4.17	6.03	3.39	4.75	4.51	4.29	4.29										
THYRO1001016	5.47	1.00	0.49	0.81	2.15	1.07	3.41	1.14	1.14										
THYRO1001023	4.57	1.75	1.46	2.49	2.37	2.27	3.16	2.69	2.69										
THYRO1001031	7	3.67	3.54	7.94	8.10	9.10	7.42	6.69	6.69*	+									
THYRO1001033	2.8	0.57	1.23	2.39	2.37	1.06	1.41	2.32	2.32										
THYRO1001062	3.82	2.25	2.08	5.76	5.14	5.15	3.45	3.99	3.99*	+									
THYRO1001063	2.69	1.60	2.09	4.12	3.13	4.17	2.95	2.51	2.51*	+									
THYRO1001071	0.69	1.53	1.22	0.98	1.08	0.43	1.16	1.21	1.21										
THYRO1001080	5.05	2.34	2.74	5.3	3.86	5.08	3.55	4.04	4.04										
THYRO1001093	3.71	2.05	1.76	6.8	6.51	4.95	3.07	3.82	3.82*	+									
THYRO1001106	2.79	1.59	1.28	2.23	2.44	2.67	1.71	3.71	3.71										
THYRO1001102	4.56	2.46	2.61	2.98	3.38	2.67	4.62	4.11	4.11										
THYRO1001104	7.28	6.54	6.58	7.94	7.41	6.48	4.57	5.35	5.35										
THYRO1001107	2.63	2.02	1.30	2.32	2.09	1.60	2.52	1.80	1.8										
THYRO1001113	1.05	0.71	0.52	0.95	1.64	0.74	2.24	3.05	3.05										

170/180

【0811】

【表340】

1817

1818

THYRO1001120	3.6	3.56	2.97	4.01	3.89	3.81	3.24	4.59	4.59				
THYRO1001121	4.68	3.13	2.03	5.64	4.07	3.90	2.70	4.05	4.05				
THYRO1001122	6.11	5.32	3.34	12.06	12.42	10.51	5.36	6.39	6.39	**	+		
THYRO1001133	6.15	4.73	4.57	9.2	11.55	7.92	6.41	7.28	7.28	*		+	
THYRO1001134	3.36	2.97	3.23	3.78	3.94	5.18	4.36	4.50	4.5		**	+	
THYRO1001142	0.74	0.74	1.04	0.72	2.52	2.41	0.96	1.79	1.79				+
THYRO1001173	15.19	9.02	12.22	26.91	29.74	31.51	28.83	31.54	31.54	**	+	**	+
THYRO1001175	1.52	0.43	1.46	2.01	0.80	2.13	0.96	1.73	1.73				
THYRO1001177	2.64	2.90	2.12	5.03	6.80	5.41	2.98	4.26	4.26	**	+		
THYRO1001189	11.01	7.39	8.79	19.93	32.38	18.70	9.07	8.97	8.97	*	+		
THYRO1001194	3.46	1.13	2.28	5.96	5.42	5.39	1.82	2.43	2.43	**	+		
THYRO1001204	4.45	2.95	2.30	6.96	6.86	8.50	3.26	4.79	4.79	**	+		
THYRO1001205	24.03	16.88	15.68	32.39	32.90	31.15	22.06	24.66	24.66	**	+		
THYRO1001213	3.76	2.34	2.06	5.73	8.42	6.51	4.19	4.49	4.49	*	+	*	+
THYRO1001224	9.88	5.89	5.95	9.43	12.54	11.82	5.58	6.76	6.76				
THYRO1001237	2.56	2.32	3.39	3.81	2.63	3.98	5.21	5.02	5.02		**	+	
THYRO1001242	27.87	23.01	22.93	21.64	25.67	32.15	25.14	28.77	28.77				
THYRO1001251	3.57	5.51	4.92	4.9	6.74	6.73	7.47	5.30	5.3				
THYRO1001262	1.72	1.10	1.83	6.36	5.01	5.41	2.24	3.79	3.79	**	+	*	+
THYRO1001266	1.55	0.64	0.79	1.26	1.48	1.18	1.70	1.12	1.12				
THYRO1001271	3.44	2.05	1.29	2.26	3.55	2.36	3.05	2.35	2.35				
THYRO1001287	3.96	1.21	1.37	3.53	2.40	2.74	3.19	2.91	2.91				
THYRO1001290	1.14	0.69	1.23	1.44	2.26	2.04	2.54	3.09	3.09	*	+	**	+
THYRO1001291	1.66	1.74	1.06	3.35	4.38	3.14	2.28	4.20	4.2	**	+	*	+
THYRO1001297	5.89	5.62	3.44	7.28	6.73	6.27	3.04	3.57	3.57				
THYRO1001302	0.7	1.17	1.36	2.14	3.01	3.14	1.40	2.26	2.26	*	+		
THYRO1001313	4.31	2.12	1.72	3.28	3.86	2.48	2.67	3.67	3.67				
THYRO1001320	4.07	2.24	2.43	7.21	7.25	7.12	3.37	4.30	4.3	**	+		
THYRO1001321	4.3	1.74	1.67	5.83	6.09	3.75	2.97	2.21	2.21				
THYRO1001322	2.79	2.55	2.39	3.89	5.05	3.82	2.48	1.98	1.98	*	+		
THYRO1001337	1.5	1.06	0.78	3.17	2.62	2.46	1.64	1.54	1.54	**	+		
THYRO1001336	5.87	4.46	7.00	13.05	17.27	14.64	6.39	6.28	6.28	**	+		
THYRO1001347	0.03	0.55	0.25	0.69	2.15	0.73	1.35	0.54	0.54				
THYRO1001358	11.06	9.93	9.25	14.71	16.38	14.53	9.85	8.62	8.62	**	+		
THYRO1001363	5.86	3.17	4.11	5.35	3.91	6.10	4.52	5.65	5.65				
THYRO1001365	5.19	2.07	3.95	4.26	3.12	4.83	2.55	3.93	3.93				
THYRO1001374	9.65	2.81	3.50	6.43	5.39	7.37	3.94	7.65	7.65				
THYRO1001401	7.01	3.08	4.71	9.44	10.37	11.91	6.83	6.19	6.19	*	+		
THYRO1001403	5.97	2.05	2.57	7.36	6.46	7.19	3.33	5.45	5.45				
THYRO1001405	5.97	3.44	4.77	7.32	6.00	9.69	6.01	5.53	5.53				
THYRO1001406	18.99	10.90	12.10	23.76	22.00	31.87	17.99	23.95	23.95	*	+		
THYRO1001411	13.78	6.66	6.31	15.64	15.28	13.18	9.35	10.33	10.33				
THYRO1001420	16.57	7.72	7.86	12.67	12.64	10.93	13.35	14.42	14.42				
THYRO1001426	12.94	7.75	6.41	21.71	18.55	24.48	11.12	13.81	13.81	**	+		
THYRO1001430	8.77	5.32	6.79	6.79	9.85	9.38	6.22	8.03	8.03				
THYRO1001434	4.36	1.78	2.26	3.34	4.87	4.04	2.27	4.13	4.13				
THYRO1001456	6.47	2.68	3.34	4.42	3.96	4.89	4.38	4.91	4.91				
THYRO1001457	6.96	3.92	4.84	6.7	6.42	9.36	5.12	7.08	7.08				
THYRO1001458	9.57	4.98	6.73	8.45	4.79	10.94	7.33	10.09	10.09				
THYRO1001459	11.09	4.54	5.24	11.67	15.21	11.24	5.79	8.41	8.41				
THYRO1001471	6.36	3.07	1.95	5.35	3.31	4.14	2.83	4.03	4.03				
THYRO1001478	6.87	2.62	2.63	3.98	3.73	5.94	4.03	6.74	6.74				
THYRO1001480	13.1	8.34	8.71	20.72	21.69	22.38	10.45	33.77	33.77	**	+		
THYRO1001481	5.7	2.94	3.90	7.61	6.97	7.30	4.44	6.00	6	*	+		
THYRO1001487	7.46	5.22	5.93	9.3	8.06	9.99	6.17	7.84	7.84	*	+		
THYRO1001495	11.89	6.81	10.31	8.41	6.19	9.91	4.61	6.76	6.76				

171/180

【0812】

【表341】

1819

1820

THYRO1001498	9.2	3.54	3.52	8.32	6.23	9.44	6.75	6.00	6										
THYRO1001510	8.51	2.92	3.62	4.12	4.26	4.21	2.96	4.74	4.74										
THYRO1001512	9.32	6.84	5.74	9.67	9.37	8.03	7.58	10.22	10.22										
THYRO1001519	9.13	4.10	4.70	9.27	7.38	9.67	6.98	8.20	8.2										
THYRO1001522	6.26	4.50	5.23	7.93	8.82	7.33	5.58	9.26	9.26	*	+								
THYRO1001523	3.53	2.10	1.99	6.46	5.54	6.24	4.04	4.29	4.29	**	+	*	+						
THYRO1001526	6.91	4.84	5.74	14.18	9.51	13.49	12.30	16.11	16.11	*	+	**	+						
THYRO1001529	2.41	1.14	1.41	2.28	1.58	4.28	2.24	2.20	2.2										
THYRO1001534	3.65	2.24	1.50	4.38	3.58	6.43	2.88	4.21	4.21										
THYRO1001537	18.2	10.50	9.67	21.59	21.38	19.81	8.19	10.14	10.14	*	+								
THYRO1001541	14.28	6.89	6.76	16.77	16.36	14.76	9.61	10.03	10.03										
THYRO1001545	3.56	2.76	2.72	3.42	3.96	4.48	3.96	4.30	4.3			*	+						
THYRO1001552	3.99	2.04	2.13	4.24	3.76	7.51	3.56	3.91	3.91										
THYRO1001563	11.96	7.39	6.70	7.96	5.68	9.41	7.19	8.07	8.07										
THYRO1001570	4.68	4.47	3.76	4.09	3.00	4.87	4.64	6.87	6.87										
THYRO1001573	8.02	5.52	6.21	6.26	3.61	8.28	6.11	6.00	6										
THYRO1001584	8.32	5.29	4.71	9.43	6.63	9.84	5.17	6.12	6.12										
THYRO1001593	2.99	0.93	1.22	3.14	4.86	2.61	2.01	4.21	4.21										
THYRO1001595	5.67	1.96	2.39	7.68	7.67	6.34	3.91	4.14	4.14	*	+								
THYRO1001596	3.89	2.66	3.80	3.78	3.65	3.11	2.98	3.57	3.57										
THYRO1001602	7.81	2.64	3.23	7.32	8.69	7.89	4.74	7.00	7										
THYRO1001605	5.26	2.56	2.24	5.13	5.05	4.87	3.48	3.41	3.41										
THYRO1001608	7.75	3.89	6.86	6.23	6.07	8.04	6.19	6.87	6.87										
THYRO1001617	14.26	9.34	10.47	17.37	15.68	19.92	9.80	12.17	12.17	*	+								
THYRO1001634	4.95	3.06	3.93	4.4	3.84	4.30	4.75	4.39	4.39										
THYRO1001637	10.18	6.14	4.65	17.45	14.38	17.46	8.06	9.17	9.17	**	+								
THYRO1001641	6.38	3.44	3.03	6.59	5.36	5.81	5.90	5.59	5.59										
THYRO1001656	4.32	2.95	2.83	3.81	4.14	7.31	4.33	5.14	5.14										
THYRO1001658	4.29	2.01	1.79	2.18	2.89	2.10	2.16	2.58	2.58										
THYRO1001661	3.1	1.45	1.64	1.98	2.33	1.46	4.01	2.50	2.5										
THYRO1001671	5.77	2.59	2.20	4.22	4.26	4.64	3.03	5.39	5.39										
THYRO1001672	6.81	4.51	5.53	5.21	5.27	6.87	6.28	6.63	6.63										
THYRO1001673	4	1.65	1.66	5.32	3.21	5.73	2.44	2.64	2.64										
THYRO1001677	6.31	4.12	3.30	6.16	7.35	6.56	2.26	3.46	3.46										
THYRO1001683	8.24	4.40	3.37	4.91	4.29	8.77	5.76	11.28	11.28										
THYRO1001700	4.49	4.00	2.73	4.05	4.60	4.19	4.01	4.47	4.47										
THYRO1001702	15.24	5.52	7.38	9.42	10.75	10.20	8.66	10.47	10.47										
THYRO1001703	9.25	6.47	6.51	7.26	6.71	8.49	10.46	8.63	8.63										
THYRO1001706	4.3	2.92	3.16	3.43	6.68	7.52	2.62	4.78	4.78	*	+								
THYRO1001721	5.23	3.35	2.76	6.77	6.22	4.74	5.26	7.04	7.04			*	+						
THYRO1001725	4.92	2.94	2.29	5.59	6.33	8.71	2.75	4.72	4.72	*	+								
THYRO1001730	24.29	13.18	13.43	14.02	17.03	13.70	21.66	22.76	22.76										
THYRO1001738	9.75	4.90	4.82	9.04	5.85	7.43	4.92	7.98	7.98										
THYRO1001743	4	3.23	1.86	3.1	3.27	3.05	4.23	2.96	2.96										
THYRO1001745	2.52	1.07	1.25	1.89	1.88	1.53	1.75	2.88	2.88										
THYRO1001746	4.33	2.26	1.61	3.33	3.91	4.02	3.18	3.68	3.68										
THYRO1001770	12.11	9.28	9.48	15.96	14.31	15.08	8.82	12.34	12.34	**	+								
THYRO1001772	5.17	2.74	2.39	6.3	7.93	7.50	3.31	3.90	3.9	*	+								
THYRO1001778	15.04	12.42	11.52	14.35	19.15	15.23	13.02	16.18	16.18										
THYRO1001793	14.69	6.79	7.43	12.47	11.55	9.29	6.11	8.93	8.93										
THYRO1001796	11.97	7.13	5.24	9.74	7.66	8.26	8.13	8.63	8.63										
THYRO1001800	6.25	3.20	2.10	5.72	6.37	4.12	5.27	7.47	7.47										
THYRO1001803	19.67	13.46	12.81	14.42	14.10	16.43	18.72	18.13	18.13										
THYRO1001809	3.63	3.46	2.67	3.26	4.76	4.46	2.91	5.54	5.54										
THYRO1001817	6.44	6.33	5.18	4.47	3.95	4.74	8.83	7.32	7.32	*		*	+						
THYRO1001819	5.55	5.75	5.06	8.24	5.95	6.83	5.79	6.86	6.86										

172/180

【0813】

【表342】

1821

1822

JHYRO1001828	5.58	5.56	4.00	9.32	9.83	9.03	4.86	6.29	6.29	**	+	
JHYRO1001854	20.22	7.97	7.27	24.83	26.41	23.02	14.19	14.50	14.5	*	+	
JHYRO1001895	4.5	1.82	1.66	2.69	3.40	3.20	2.51	2.17	2.17			
JHYRO1001907	6.37	2.87	2.77	7.43	8.35	6.14	3.08	4.67	4.67			
TRACH1000006	1.82	2.19	1.60	2.9	3.42	2.53	2.58	3.05	3.05	*	+	+
TRACH1000013	2.15	1.13	1.31	1.45	1.80	3.25	1.50	1.76	1.76			
TRACH1000074	3.42	3.57	4.39	5.62	7.83	7.88	4.19	10.27	10.27	*	+	
TRACH1000095	2.45	2.91	2.44	3.1	3.04	4.04	2.50	2.45	2.45			
TRACH1000103	7.43	5.84	4.56	10.07	11.80	13.53	5.10	8.65	8.65	*	+	
TRACH1000108	3.15	1.08	0.60	4.55	2.50	3.75	3.10	1.49	1.49			
TRACH1000126	6.59	4.83	4.15	6.73	6.75	6.24	2.66	4.52	4.52			
TRACH1000146	4.1	2.48	3.17	3.77	4.50	3.73	2.81	3.85	3.85			
TRACH1000168	2.88	1.73	0.69	2.15	3.29	1.84	1.31	2.46	2.46			
TRACH1000184	9.18	5.15	6.68	9.87	12.29	12.18	7.92	7.13	7.13	*	+	
VESEN1000004	1.43	3.20	2.03	4.77	4.23	4.76	2.44	2.90	2.9	*	+	
VESEN1000007	4.67	3.71	3.03	4.92	4.79	4.78	3.45	3.27	3.27			
VESEN1000013	3.8	4.40	3.49	6.08	5.11	8.39	4.08	5.78	5.78			
VESEN1000028	10.32	4.13	4.71	9.23	9.35	9.07	7.29	12.27	12.27			
VESEN1000049	7.75	3.60	4.26	7.63	6.94	7.73	4.60	5.95	5.95			
VESEN1000100	14.3	7.29	8.52	11.77	17.29	16.55	10.06	12.85	12.85			
VESEN1000107	8.09	2.86	4.55	5.28	4.93	5.96	5.50	6.28	6.28			
VESEN1000117	4.56	2.53	3.13	3.83	3.21	3.98	3.40	4.83	4.83			
VESEN1000122	6	2.68	4.24	3.89	4.52	7.18	4.38	7.65	7.65			
VESEN1000137	2.93	1.73	1.82	1.57	3.65	3.17	2.10	3.43	3.43			
VESEN1000195	14.98	5.35	5.89	8.11	8.22	6.74	10.54	12.97	12.97			
VESEN1000215	2.26	0.13	1.20	1.57	1.68	0.85	0.67	1.63	1.63			
VESEN1000279	26.58	15.13	14.91	21.43	14.13	23.59	19.30	20.07	20.07			
VESEN1000363	15.34	8.73	10.79	17.48	16.61	12.88	9.72	13.31	13.31			
VESEN1000388	9.91	6.40	6.52	7.89	4.01	10.40	6.86	10.14	10.14			
VESEN1000394	12.12	6.72	8.23	12.56	8.96	9.43	5.04	9.23	9.23			
VESEN1000410	10.78	2.59	2.39	6.85	3.24	4.07	5.06	8.94	8.94			
VESEN1000411	6.18	3.27	4.03	5.74	3.11	6.71	4.21	5.31	5.31			
VESEN1000415	9.24	6.34	4.20	8.16	6.27	5.95	4.08	7.14	7.14			
VESEN1000440	9.05	5.57	4.80	8.89	8.64	8.72	5.45	8.25	8.25			
VESEN1000452	7.8	4.72	5.60	4.86	5.38	4.21	6.76	5.77	5.77			
VESEN1000539	346.75	188.95	244.65	158.74	166.73	144.68	64.90	151.18	151.2			
VESEN1000544	4.46	3.39	3.95	4.07	2.23	3.58	2.95	2.93	2.93	*	-	
VESEN1000557	6.06	4.00	4.41	6.38	3.08	5.06	6.10	7.77	7.77	*	+	
VESEN1000575	7.82	4.18	4.70	6.03	4.15	4.58	5.87	6.64	6.64			
VESEN1000585	9.14	4.16	5.29	6.86	6.14	7.55	4.21	6.93	6.93			
VESEN1000592	1.51	0.34	0.06	1.48	0.81	0.75	1.11	0.98	0.98			
VESEN1000658	9.42	5.35	3.63	6.6	8.13	5.18	7.65	9.88	9.88			
VESEN1000669	30.52	16.02	17.70	27.74	22.51	23.12	18.76	27.04	27.04			
VESEN1000743	12.62	7.52	8.72	9.64	10.40	10.72	6.57	9.41	9.41			
VESEN1000752	31.33	20.56	19.92	44.49	19.58	40.73	21.19	32.70	32.7			
VESEN1000761	23.86	13.01	17.50	12.45	9.94	17.39	8.43	10.21	10.21			
VESEN2000039	77.69	44.95	56.28	57.5	43.19	64.97	60.33	69.54	69.54			
VESEN2000102	7.33	4.99	5.35	6.83	4.25	7.08	6.69	8.37	8.37			
VESEN2000164	5.18	3.46	3.31	9.13	9.21	6.82	3.36	3.89	3.89	*	+	
VESEN2000175	1.73	0.97	0.12	1.01	1.92	1.13	0.88	1.17	1.17			
VESEN2000186	19.39	12.37	11.60	17.79	16.80	18.96	15.99	20.01	20.01			
VESEN2000199	28.49	19.51	19.01	18.68	21.47	33.21	23.58	23.01	23.01			
VESEN2000200	6.32	1.63	3.02	5.06	3.00	3.70	3.04	4.39	4.39			
VESEN2000204	4.52	1.87	3.26	2.47	1.87	2.02	2.17	3.09	3.09			
VESEN2000218	6.43	3.74	5.10	6.59	6.27	8.76	4.84	5.35	5.35			
VESEN2000230	5.26	2.88	3.63	6.04	5.20	6.82	6.20	5.85	5.85	*	+	

173/180

【0814】

【表343】

1823

1824

VESEN2000272	6.36	2.52	3.61	13.68	15.50	9.23	6.37	6.11	6.11	*		
VESEN2000292	5.8	3.32	3.03	6.33	5.54	5.31	4.11	3.82	3.82			
VESEN2000323	3.64	2.70	3.46	7.25	6.60	6.63	4.13	6.99	6.99	**	+	+
VESEN2000377	16.91	9.24	9.32	14.89	11.98	16.05	16.51	12.53	12.53			
VESEN2000378	3.41	1.69	2.05	2.7	1.99	2.52	3.68	4.21	4.21		*	+
VESEN2000330	9.06	4.24	3.98	4	3.94	4.40	7.56	5.58	5.58			
VESEN2000336	3.29	2.35	2.63	3.19	2.56	2.84	2.06	2.38	2.38			
VESEN2000354	8.7	4.46	4.22	7.46	6.89	5.83	5.63	5.02	5.02			
VESEN2000378	3.42	2.15	2.25	4.13	2.42	1.91	1.92	2.61	2.61			
VESEN2000379	11.63	7.79	4.82	10.74	10.07	12.49	7.29	10.70	10.7			
VESEN2000397	3.37	1.29	1.36	2.39	2.24	1.99	1.18	3.19	3.19			
VESEN2000416	3.83	2.34	1.55	2.15	2.33	2.91	2.47	2.28	2.28			
VESEN2000430	2.88	0.98	1.36	1.52	0.23	0.52	0.64	1.63	1.63			
VESEN2000430	2.62	1.65	1.71	1.89	2.49	1.83	0.78	2.97	2.97			
VESEN2000448	2.86	2.67	1.17	1	2.01	2.37	2.33	2.73	2.73			
VESEN2000449	8.25	5.92	4.67	9.14	8.56	10.89	3.16	6.55	6.55			
VESEN2000456	5.37	3.06	1.86	3.12	2.41	3.57	2.05	2.65	2.65			
VESEN2000562	7.78	4.41	5.30	5.84	5.51	4.92	4.30	6.44	6.44			
VESEN2000573	0.6	0.35	0.41	0.67	0.40	0.67	1.28	2.60	2.6		*	+
VESEN2000604	5.64	1.48	1.85	3.25	2.37	2.19	2.91	4.05	4.05			
VESEN2000614	25.21	13.24	16.03	20.97	19.46	20.96	23.97	21.61	21.61			
VESEN2000638	1.7	1.28	1.62	1.56	1.85	1.20	2.41	1.35	1.35			
VESEN2000641	1.73	2.11	1.08	1.79	1.66	1.77	1.14	1.95	1.95			
VESEN2000645	3.09	2.77	2.30	2.12	2.14	1.71	1.70	3.15	3.15			
Y79AA1000013	10.79	7.40	5.68	11.91	9.74	8.63	7.82	6.74	6.74			
Y79AA1000030	13.95	8.47	8.24	10.96	9.10	13.62	9.47	12.29	12.29			
Y79AA1000033	16.96	12.16	9.55	7.65	10.20	8.44	7.18	10.76	10.76			
Y79AA1000037	2.11	1.49	0.71	2.23	2.21	3.27	2.75	2.51	2.51		*	+
Y79AA1000041	2.2	2.48	1.77	2.69	2.36	2.74	2.02	3.82	3.82			
Y79AA1000059	7.6	6.90	6.65	10.99	11.69	12.90	4.30	7.70	7.7	**	+	
Y79AA1000065	22.39	17.36	15.96	24.43	21.67	25.09	14.43	16.06	16.06			
Y79AA1000081	42.69	41.35	51.24	111.38	113.45	103.25	45.62	16.30	16.3	**	+	
Y79AA1000127	22.29	16.01	11.79	12.57	10.65	7.07	3.98	5.58	5.58		*	+
Y79AA1000130	6.17	3.27	2.89	10.01	8.60	9.63	4.89	5.13	5.13	**	+	
Y79AA1000131	448.19	235.19	299.39	399.75	486.28	438.12	277.73	304.61	304.6			
Y79AA1000134	8.96	7.49	5.25	6.6	6.53	6.62	9.23	10.69	10.69			
Y79AA1000143	9.99	4.29	8.06	7.58	8.06	8.95	6.96	8.30	8.3			
Y79AA1000144	8.55	7.18	6.04	6.31	5.55	6.00	4.05	4.40	4.4		*	-
Y79AA1000150	18.22	14.18	15.26	14.89	15.33	21.06	9.92	9.91	9.91	**		
Y79AA1000153	201.67	139.66	172.85	190.71	189.25	179.30	103.81	119.17	119.2	*		
Y79AA1000166	6.51	3.61	2.42	6.7	8.84	4.48	3.56	4.21	4.21			
Y79AA1000179	15.16	9.65	7.92	10.53	9.30	7.94	4.29	5.64	5.64			
Y79AA1000181	10.66	5.30	5.63	7.26	8.22	5.85	3.94	5.98	5.98			
Y79AA1000202	18.5	15.06	12.86	18.25	18.73	23.11	15.84	25.98	25.98			
Y79AA1000207	5.87	4.02	4.27	14.67	14.22	14.10	7.48	5.85	5.85	**	+	
Y79AA1000214	29.22	23.27	20.29	36.32	37.06	45.40	22.86	25.86	25.86	*	+	
Y79AA1000222	12.84	9.84	10.93	9.21	6.89	9.06	5.29	5.66	5.66		**	
Y79AA1000226	5.63	6.09	5.68	7.41	7.20	8.09	8.84	8.79	8.79	**	+	***
Y79AA1000227	17.27	10.05	8.43	12.69	17.80	12.32	9.20	10.19	10.19			
Y79AA1000230	6.42	4.02	2.20	3.72	4.88	2.48	3.03	2.90	2.9			
Y79AA1000231	34.72	21.74	21.36	20.87	19.10	17.13	9.21	15.10	15.1			
Y79AA1000239	15.79	9.79	7.30	10.27	13.40	11.55	12.77	13.82	13.82			
Y79AA1000258	4.05	3.20	3.26	4.22	5.80	4.84	3.99	4.25	4.25			
Y79AA1000268	7.27	4.70	4.79	10.11	6.83	6.96	5.20	6.24	6.24			
Y79AA1000269	3.42	2.81	2.55	4.54	6.08	5.88	5.38	5.60	5.6	**	+	***
Y79AA1000270	3.64	4.17	2.51	5.74	6.14	5.66	3.62	4.41	4.41	**	+	+

174/180

1825

1826

Y79AA1000288	11.25	5.37	6.77	11.8	13.92	12.66	5.46	9.54	9.54				
Y79AA1000285	4.46	1.52	2.70	3.31	1.78	2.60	2.43	3.53	3.53				
Y79AA1000292	3.61	2.65	3.31	10.15	10.34	10.77	4.41	5.66	5.66	**	+	+	+
Y79AA1000307	12.46	9.65	13.13	11.87	8.54	13.75	5.29	6.68	6.68			**	-
Y79AA1000313	15.46	6.94	8.62	10.28	12.44	14.87	10.41	13.90	13.9				
Y79AA1000314	14.81	9.18	10.30	22.74	18.92	27.80	24.11	31.46	31.46	*	+	**	+
Y79AA1000328	3.09	1.87	2.24	2.09	2.55	2.73	1.78	2.96	2.96				
Y79AA1000334	7.09	3.70	2.56	5.55	4.48	4.69	3.41	4.25	4.25				
Y79AA1000342	35.87	15.66	15.62	22.36	17.70	23.91	21.00	29.07	29.07				
Y79AA1000346	17.41	15.57	12.74	9.41	9.10	10.71	4.23	5.49	5.49	*	-	**	-
Y79AA1000347	23.11	14.24	15.07	23.5	39.38	38.47	19.81	25.73	25.73	*	+		
Y79AA1000349	19.76	10.53	12.68	20.31	16.01	21.05	12.82	17.27	17.27				
Y79AA1000355	4.87	2.42	3.06	7.28	6.44	8.31	4.76	6.17	6.17	*	+		
Y79AA1000369	6.76	2.87	3.15	4.62	3.69	3.41	4.31	4.40	4.4				
Y79AA1000388	25.23	15.44	16.71	26.79	21.25	29.10	12.60	17.85	17.85				
Y79AA1000392	14.91	8.34	9.71	13.34	7.02	19.13	9.61	11.82	11.82				
Y79AA1000405	24.03	14.82	7.15	15.39	22.71	12.76	14.12	17.35	17.35				
Y79AA1000410	24.25	16.23	12.97	37.19	36.14	36.35	20.62	22.06	22.06	**	+		
Y79AA1000420	1.83	1.06	1.88	2.33	1.74	3.81	1.85	2.84	2.84				
Y79AA1000423	7.25	4.11	5.48	9.75	7.86	8.44	5.00	5.45	5.45	*	+		
Y79AA1000426	5.29	3.84	5.55	4.45	2.88	4.33	3.32	3.94	3.94				
Y79AA1000432	3.27	2.71	3.28	1.62	1.68	2.55	1.63	2.22	2.22	*	-	*	-
Y79AA1000453	141.24	53.68	107.37	81.71	59.38	81.50	30.05	43.77	43.77				
Y79AA1000465	3.59	1.59	2.02	2.43	1.32	2.55	1.25	3.10	3.1				
Y79AA1000469	14.01	11.65	7.90	12.08	10.53	7.10	8.31	7.33	7.33				
Y79AA1000480	4.69	1.58	1.60	4.05	2.82	2.60	2.60	2.44	2.44				
Y79AA1000502	12.81	5.39	8.31	9.83	13.49	9.32	5.96	11.12	11.12				
Y79AA1000511	6.28	4.42	6.32	6.26	4.77	4.40	6.38	6.79	6.79				
Y79AA1000534	17.26	8.63	8.69	10.74	7.23	7.43	4.39	5.56	5.56				
Y79AA1000538	6.63	3.28	4.52	10.32	7.26	8.06	5.36	6.47	6.47	*	+		
Y79AA1000539	19.25	8.27	12.78	24.31	26.47	21.68	9.27	11.72	11.72	*	+		
Y79AA1000540	11.13	5.92	6.15	9.13	9.09	8.44	6.65	9.21	9.21				
Y79AA1000560	173.06	134.34	94.53	202.66	161.69	169.55	95.78	139.04	139				
Y79AA1000574	2.89	2.45	2.28	4.12	2.97	2.60	1.96	2.63	2.63				
Y79AA1000584	3.2	1.68	1.63	1.75	2.10	2.56	2.05	2.41	2.41				
Y79AA1000589	8.66	5.80	5.36	6.79	3.71	6.73	6.49	7.62	7.62				
Y79AA1000598	5.98	2.97	4.18	3.57	3.29	6.10	4.35	4.63	4.63				
Y79AA1000600	6.57	3.44	3.89	3.3	2.25	3.48	2.55	2.77	2.77				
Y79AA1000609	6.92	3.42	2.75	2.76	4.04	6.09	4.13	5.52	5.52				
Y79AA1000618	58.41	20.55	40.08	29.91	37.02	38.12	11.43	14.49	14.49	*	-		
Y79AA1000627	6.08	3.22	3.45	5.69	5.50	4.18	4.40	3.93	3.93				
Y79AA1000636	38.19	23.55	23.75	16.84	22.87	15.14	9.44	11.05	11.05	*	-		
Y79AA1000649	8.69	4.34	4.67	4.61	4.61	4.01	3.93	8.79	8.79				
Y79AA1000654	5.76	3.08	3.22	5.58	4.90	5.16	3.04	4.23	4.23				
Y79AA1000673	5.03	2.72	1.36	3.23	1.94	2.41	3.39	4.06	4.06				
Y79AA1000674	10.61	7.11	11.17	10.18	8.67	10.62	6.76	10.00	10				
Y79AA1000678	7.25	4.89	6.06	10.19	6.81	7.33	5.06	5.92	5.92				
Y79AA1000682	24.87	16.17	18.30	22.46	26.14	12.58	16.87	20.51	20.51				
Y79AA1000683	15.32	7.96	8.21	6.64	7.38	6.63	4.88	6.13	6.13				
Y79AA1000697	54.8	30.85	37.16	42.84	41.24	37.90	36.31	42.61	42.61				
Y79AA1000700	9.78	3.97	5.64	3.6	3.66	3.51	4.90	7.29	7.29				
Y79AA1000702	17.82	9.90	9.05	9.33	10.05	8.49	5.94	9.28	9.28				
Y79AA1000704	2.05	0.88	1.35	0.9	1.80	0.91	1.41	1.66	1.66				
Y79AA1000705	2.45	1.63	1.24	3.99	2.73	3.50	2.26	2.26	2.26	*	+		
Y79AA1000717	11.47	6.51	7.68	14.26	8.53	10.93	5.99	11.28	11.28				
Y79AA1000723	6.59	5.15	4.02	3.83	3.18	4.48	1.26	1.65	1.65	**	-		

175/180

【0816】

【表345】

Y79AA1000724	28.17	13.18	13.80	13.88	13.98	11.98	3.06	4.28	4.28		*	-
Y79AA1000726	8.11	5.46	4.24	6.09	4.77	4.52	5.43	7.82	7.82			
Y79AA1000734	3.88	2.62	2.34	5.17	3.55	4.31	2.92	6.05	6.05			
Y79AA1000748	3.95	1.81	1.83	2.64	2.02	2.92	1.57	2.24	2.24			
Y79AA1000750	10.39	6.10	4.86	9.81	8.59	9.78	5.43	7.43	7.43			
Y79AA1000752	2.87	0.53	1.08	2.54	2.81	2.11	1.32	1.59	1.59			
Y79AA1000774	5.72	4.59	2.86	2.14	2.79	5.77	3.53	3.76	3.76			
Y79AA1000776	4.35	4.36	2.86	3.71	4.12	5.01	3.48	3.90	3.9			
Y79AA1000777	11.76	6.21	5.54	6.56	11.90	10.17	6.16	6.66	6.66			
Y79AA1000778	13.22	6.87	8.41	14.77	13.90	13.40	7.19	13.72	13.72			
Y79AA1000782	7.86	4.93	5.51	5.52	4.90	5.05	5.46	7.23	7.23			
Y79AA1000784	12.43	9.12	11.59	13	14.52	14.46	11.05	11.31	11.31			
Y79AA1000794	4.35	2.95	2.89	4.43	4.95	3.90	3.24	3.10	3.1			
Y79AA1000800	2.57	2.36	2.08	3	3.32	3.30	2.93	3.69	3.69	**	+	+
Y79AA1000802	1.85	1.48	1.65	1	0.76	1.64	0.34	1.23	1.23			
Y79AA1000805	4.24	3.55	2.28	2.22	3.19	3.89	2.71	4.15	4.15			
Y79AA1000814	14.61	9.83	7.28	9.51	9.83	6.77	3.86	4.30	4.3		*	-
Y79AA1000823	12.6	9.53	9.56	15.44	14.21	12.23	9.08	15.12	15.12			
Y79AA1000824	4.44	3.44	2.16	2.49	3.58	2.72	2.72	3.74	3.74			
Y79AA1000827	3.1	1.46	1.84	2.99	1.29	1.77	1.89	2.61	2.61			
Y79AA1000831	5.49	4.85	5.37	3.74	4.89	3.85	3.76	5.38	5.38			
Y79AA1000833	40.22	31.45	37.17	40.96	46.51	50.53	34.20	40.04	40.04			
Y79AA1000850	2.09	2.81	2.57	4.27	3.76	4.02	3.33	2.26	2.26	**	+	
Y79AA1000856	6.74	5.50	6.27	7.85	6.17	10.60	4.73	5.48	5.48			
Y79AA1000862	12.52	7.78	4.39	13.89	9.86	8.13	7.63	7.94	7.94			
Y79AA1000876	8.46	4.16	4.01	6.87	6.89	6.26	3.75	5.07	5.07			
Y79AA1000888	1.47	1.34	1.40	1.56	1.46	1.29	1.98	1.99	1.99	**	+	
Y79AA1000902	16.38	10.81	14.11	11.4	9.46	11.97	5.88	7.23	7.23		*	-
Y79AA1000935	16.25	11.98	13.09	25.37	21.17	25.92	23.44	29.28	29.28	**	+	**
Y79AA1000959	3.1	2.66	3.26	3.18	3.69	2.84	2.68	4.50	4.5			
Y79AA1000963	1.8	2.34	1.77	4.45	3.80	4.94	2.33	2.34	2.34	**	+	
Y79AA1000963	43.49	20.23	23.14	40.9	40.35	45.98	17.97	19.24	19.24			
Y79AA1000966	8	6.62	3.05	7.53	7.08	4.56	6.48	5.59	5.59			
Y79AA1000967	11.14	8.37	5.21	15.29	15.02	10.80	8.86	10.67	10.67			
Y79AA1000968	11.05	6.63	3.78	6.52	9.03	6.81	4.66	7.08	7.08			
Y79AA1000969	4.13	3.63	3.19	4.09	3.12	3.96	2.88	4.11	4.11			
Y79AA1000976	2.07	1.66	1.63	2.46	2.43	2.76	2.15	3.14	3.14	*	+	+
Y79AA1000978	3.15	2.68	2.59	3.19	2.43	2.99	1.56	2.57	2.57			
Y79AA1000985	4.53	6.21	3.11	9.92	6.66	7.93	4.84	4.19	4.19			
Y79AA1000989	27.14	18.46	21.17	22.61	22.40	25.64	17.86	17.83	17.83			
Y79AA1000991	14.41	7.65	8.70	14.5	16.91	8.11	10.68	10.04	10.04			
Y79AA1001013	35.7	19.64	14.11	24.63	29.38	32.01	18.46	27.65	27.65			
Y79AA1001014	8.41	5.13	3.58	6.96	7.27	8.35	6.51	8.47	8.47			
Y79AA1001019	6.41	3.32	4.05	4.98	4.88	5.75	4.58	5.04	5.04			
Y79AA1001020	13.26	4.81	6.74	9.29	9.05	11.19	6.66	10.83	10.83			
Y79AA1001023	3.99	2.27	3.29	3.71	4.41	3.42	4.24	3.90	3.9			
Y79AA1001030	4.36	2.82	3.64	7.73	6.53	9.26	7.69	8.68	8.68	**	+	**
Y79AA1001035	-0.01	7.25	7.50	9.11	6.84	10.21	7.88	15.95	15.95			
Y79AA1001041	8.33	4.39	3.51	5.69	4.65	4.21	2.70	5.79	5.79			
Y79AA1001043	18.02	12.74	10.03	8.74	8.44	9.66	9.73	11.39	11.39			
Y79AA1001048	5.98	4.37	5.02	5.57	4.35	5.46	5.24	5.86	5.86			
Y79AA1001056	2.8	1.67	2.69	4.83	3.64	3.93	2.91	3.52	3.52	*	+	
Y79AA1001061	4.66	2.07	2.99	8.42	5.16	8.18	3.38	5.08	5.08	*	+	
Y79AA1001062	4.59	1.72	3.28	8.74	7.23	9.33	3.92	5.55	5.55	**	+	
Y79AA1001068	7.33	4.55	5.57	10.85	9.20	12.48	5.68	6.64	6.64	*	+	
Y79AA1001073	12.4	6.75	7.01	7.75	5.93	9.79	5.80	7.72	7.72			

1830

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177/180

1831

1832

Y79AA1001548	10.61	7.08	4.15	16.42	14.68	15.82	9.30	9.38	9.38	*	+		
Y79AA1001555	7.52	5.37	3.80	6.53	5.95	5.70	7.04	7.00	7				
Y79AA1001562	13.12	10.40	12.01	18.73	17.97	15.42	12.97	18.83	18.83	*	+		
Y79AA1001581	2.59	2.12	1.33	2.27	2.33	1.95	1.31	2.40	2.4				
Y79AA1001585	1.89	1.52	2.52	3.13	3.14	3.51	2.68	3.89	3.89	*	+	*	+
Y79AA1001592	8.75	5.76	6.22	9.06	9.03	12.16	6.95	10.71	10.71				
Y79AA1001594	2.44	2.99	2.99	4.89	6.76	6.84	2.08	3.52	3.52	**	+		
Y79AA1001603	41.01	29.22	27.39	35.33	47.15	41.79	19.68	22.24	22.24				
Y79AA1001613	11.06	8.37	6.50	10.25	10.82	7.55	6.69	6.52	6.52				
Y79AA1001630	0.95	0.54	0.85	1.19	0.72	0.95	1.19	0.88	0.88				
Y79AA1001647	6.2	2.96	3.68	2.82	5.76	5.40	3.17	4.07	4.07				
Y79AA1001664	13.83	6.76	7.31	10.57	12.90	8.91	7.51	7.68	7.68				
Y79AA1001665	3.6	3.81	4.37	4.15	4.52	5.51	3.17	4.23	4.23				
Y79AA1001679	14	9.57	9.87	11.81	14.25	13.41	7.94	7.63	7.63				
Y79AA1001692	3.06	2.79	3.66	3.62	3.64	6.60	2.78	2.76	2.76				
Y79AA1001696	0.47	0.94	0.29	1.8	1.18	2.00	1.48	1.81	1.81	*	+	**	+
Y79AA1001705	5.59	4.16	3.52	5.12	5.14	5.00	3.05	4.02	4.02				
Y79AA1001711	17.19	10.51	9.53	37.34	40.06	24.12	26.85	27.39	27.39	*	+	**	+
Y79AA1001717	1.38	0.95	0.69	2.28	1.17	1.95	0.86	2.01	2.01				
Y79AA1001719	3.1	2.90	1.65	4.96	4.48	2.69	2.06	2.48	2.48				
Y79AA1001727	5.47	4.87	4.29	8.17	8.05	7.12	4.94	6.45	6.45	**	+		
Y79AA1001750	20.76	27.54	23.83	38.95	38.37	32.83	22.83	25.62	25.62	*	+		
Y79AA1001760	6.22	6.83	3.78	10.14	8.09	8.51	8.09	4.11	4.11	*	+		
Y79AA1001777	4.19	4.98	4.30	10.69	9.61	8.63	5.89	5.49	5.49	**	+	*	+
Y79AA1001781	1.41	(0.02)	0.49	0.49	0.41	1.88	0.28	0.56	0.56				
Y79AA1001787	6.73	4.26	4.09	6.64	5.23	7.45	4.25	5.24	5.24				
Y79AA1001793	7.3	4.12	4.31	5.83	5.04	3.68	5.12	4.48	4.48				
Y79AA1001795	3	0.80	2.09	2.69	3.85	3.29	1.73	3.18	3.18				
Y79AA1001799	5.26	2.91	2.67	5.21	5.65	6.10	3.13	5.77	5.77				
Y79AA1001800	4.16	2.57	3.82	5.16	2.55	3.90	3.53	6.79	6.79				
Y79AA1001801	6.56	3.89	3.46	8.87	3.49	7.02	3.18	4.68	4.68				
Y79AA1001803	6.72	4.12	3.95	5.51	7.22	5.68	5.48	5.55	5.55				
Y79AA1001805	22.35	9.91	10.35	15.2	27.86	21.20	9.25	13.14	13.14				
Y79AA1001807	6.96	2.99	4.40	6.3	4.51	3.72	4.95	5.25	5.25				
Y79AA1001827	8.38	3.69	5.67	7.55	7.81	11.23	9.11	12.46	12.46			*	+
Y79AA1001846	4.43	2.15	3.75	6.2	4.92	5.41	3.96	7.82	7.82				
Y79AA1001848	2.85	1.48	2.40	3.01	2.43	2.61	2.57	2.46	2.46				
Y79AA1001853	13.89	10.72	11.89	14.4	8.43	13.46	12.95	13.31	13.31				
Y79AA1001863	15.14	7.58	9.41	15.02	11.89	14.02	8.02	12.33	12.33				
Y79AA1001866	9.57	4.75	5.85	11.97	24.49	9.54	5.28	9.21	9.21				
Y79AA1001874	1.66	0.73	0.26	0.48	1.10	0.61	0.67	0.63	0.63				
Y79AA1001875	9	6.56	7.74	8.02	10.17	8.54	9.22	11.36	11.36			*	+
Y79AA1001907	117.42	47.16	76.24	98.59	98.47	94.40	33.03	51.77	51.77				
Y79AA1001908	2.02	0.84	1.62	1.52	1.88	1.08	1.03	1.18	1.18				
Y79AA1001923	4.54	1.74	1.64	1.87	1.96	1.62	3.56	1.90	1.9				
Y79AA1001927	7.1	4.39	6.61	6.81	4.65	6.65	7.02	7.63	7.63				
Y79AA1001930	11.14	5.72	8.19	8.38	8.79	9.40	6.07	5.53	5.53				
Y79AA1001932	4.55	2.74	2.35	3.75	4.07	2.57	2.61	2.55	2.55				
Y79AA1001933	5.44	2.77	4.71	4.94	5.23	2.92	4.54	4.08	4.08				
Y79AA1001942	5.27	2.57	3.54	3.47	2.89	2.27	3.23	4.03	4.03				
Y79AA1001963	16.6	6.83	11.89	12.17	15.10	11.74	8.04	11.18	11.18				
Y79AA1001968	19.06	9.81	14.73	19.14	20.00	14.84	13.89	19.22	19.22				
Y79AA1001983	8.12	2.93	5.67	4.13	3.77	4.79	3.53	4.78	4.78				
Y79AA1002000	8.2	3.32	3.60	7.79	5.70	5.11	5.48	4.05	4.05				
Y79AA1002084	31.21	14.32	20.67	18.62	19.17	22.54	12.67	21.19	21.19				
Y79AA1002088	6.53	5.69	5.73	9.64	7.46	8.70	4.99	6.08	6.08	*	+		

178/180

1833

1834

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179/180

【0820】

* * 【表 3 4 9】

Y79AA10023461	5.46	3.35	2.57	6.5	7.83	6.14	2.75	4.60	4.6	*	+		
Y79AA1002345	1.93	1.66	1.86	2.93	2.21	2.54	1.34	2.05	2.05	*	+		
Y79AA10023473	3.38	1.43	1.37	3.37	2.29	2.38	2.95	2.21	2.21				
Y79AA10023476	434.81	300.04	466.40	120.28	171.61	120.00	316.81	454.58	454.6	**	-		
Y79AA10023478	5.45	6.92	5.32	7.99	10.13	8.03	4.87	4.92	4.92	*	+		
Y79AA10023481	11.63	11.08	9.56	16.28	16.98	14.53	7.89	7.01	7.01	**	+	++	
Y79AA10023488	4.34	4.47	7.01	11.41	12.79	9.45	5.70	6.37	6.37	*	+		
Y79AA10023499	4.43	1.48	1.47	4.2	2.82	2.25	3.39	3.35	3.35				
Y79AA1002447	1.81	1.09	1.32	2.36	2.58	2.43	1.55	2.35	2.35	**	+		
Y79AA1002413	15.88	6.76	10.60	19.95	26.46	17.33	9.58	12.56	12.56				
Y79AA1002416	5.12	2.89	2.97	4.45	4.32	5.10	4.13	4.19	4.19				
Y79AA1002439	2.82	1.17	1.77	2.75	1.85	2.91	4.10	5.62	5.62			+	+
Y79AA1002431	4.04	2.82	3.86	2.53	3.84	4.86	4.06	5.56	5.56				
Y79AA1002433	11.76	5.78	6.28	9.49	4.53	7.78	4.34	8.17	8.17				
Y79AA1002445	10.95	9.11	9.11	11.15	8.78	14.80	10.37	11.14	11.14				
Y79AA1002461	10.04	5.58	4.92	9.55	8.99	8.05	5.89	7.75	7.75				
Y79AA1002466	22.18	13.94	11.33	23.59	18.02	25.25	10.79	17.76	17.76				
Y79AA1002471	5.76	3.00	5.65	6.94	8.49	9.26	5.31	7.89	7.89	*	+		
Y79AA1002472	12.12	5.83	9.20	16.86	14.60	20.34	6.74	12.38	12.38	*	+		
Y79AA1002474	3.46	0.84	1.92	1.74	1.49	1.64	2.77	1.35	1.35				
Y79AA1002482	13.92	8.55	11.10	23.82	23.90	29.62	10.40	14.99	14.99	**	+		
Y79AA1002487	1.72	0.87	1.11	1.3	1.59	1.75	1.57	1.93	1.93				
Y79AA1002490	13.58	4.80	6.45	5.13	6.72	3.78	4.31	7.19	7.19				
Y79AA1002493	5.77	2.96	3.11	8.04	10.37	7.90	4.77	5.75	5.75	*	+		
ZBVC61006378	1.43	0.95	1.01	1.16	2.05	0.47	1.35	2.06	2.06				

180/180

【0821】以下に、本発明によるcDNAの全長塩基配列 *す。

と、推定アミノ酸配列の、クローン番号との対応を示 * 【表350】

クローン名	全長塩基配列 配列名	全長塩基配列 配列番号	推定アミノ酸配列 配列番号
HEMBA1000005	C-HEMBA1000005	10468	10469
HEMBA1000030	C-HEMBA1000030	10470	
HEMBA1000046	C-HEMBA1000046	10471	
HEMBA1000050	C-HEMBA1000050	10472	
HEMBA1000076	C-HEMBA1000076	10473	10474
HEMBA1000156	C-HEMBA1000156	10475	10476
HEMBA1000158	C-HEMBA1000158	10477	10478
HEMBA1000168	C-HEMBA1000168	10479	10480
HEMBA1000185	C-HEMBA1000185	10481	10482
HEMBA1000193	C-HEMBA1000193	10483	10484
HEMBA1000227	C-HEMBA1000227	10485	10486
HEMBA1000288	C-HEMBA1000288	10487	
HEMBA1000302	C-HEMBA1000302	10488	
HEMBA1000304	C-HEMBA1000304	10489	10490
HEMBA1000307	C-HEMBA1000307	10491	10492
HEMBA1000369	C-HEMBA1000369	10493	10494
HEMBA1000387	C-HEMBA1000387	10495	
HEMBA1000392	C-HEMBA1000392	10496	
HEMBA1000460	C-HEMBA1000460	10497	
HEMBA1000488	C-HEMBA1000488	10498	10499
HEMBA1000491	C-HEMBA1000491	10500	10501
HEMBA1000501	C-HEMBA1000501	10502	
HEMBA1000508	C-HEMBA1000508	10503	
HEMBA1000520	C-HEMBA1000520	10504	
HEMBA1000531	C-HEMBA1000531	10505	10506
HEMBA1000534	C-HEMBA1000534	10507	
HEMBA1000555	C-HEMBA1000555	10508	10509
HEMBA1000568	C-HEMBA1000568	10510	
HEMBA1000588	C-HEMBA1000588	10511	
HEMBA1000608	C-HEMBA1000608	10512	10513
HEMBA1000636	C-HEMBA1000636	10514	10515
HEMBA1000682	C-HEMBA1000682	10516	
HEMBA1000686	C-HEMBA1000686	10517	10518
HEMBA1000719	C-HEMBA1000719	10519	10520
HEMBA1000727	C-HEMBA1000727	10521	10522
HEMBA1000752	C-HEMBA1000752	10523	
HEMBA1000817	C-HEMBA1000817	10524	10525
HEMBA1000851	C-HEMBA1000851	10526	10527
HEMBA1000867	C-HEMBA1000867	10528	
HEMBA1000869	C-HEMBA1000869	10529	
HEMBA1000872	C-HEMBA1000872	10530	10531
HEMBA1000910	C-HEMBA1000910	10532	10533
HEMBA1000918	C-HEMBA1000918	10534	
HEMBA1000919	C-HEMBA1000919	10535	10536

(920)

特開 2002-191363

1837

1838

HEMBA1000946	C-HEMBA1000946	10537	10538
HEMBA1000968	C-HEMBA1000968	10539	
HEMBA1000971	C-HEMBA1000971	10540	10541
HEMBA1000975	C-HEMBA1000975	10542	
HEMBA1001009	C-HEMBA1001009	10543	10544
HEMBA1001022	C-HEMBA1001022	10545	
HEMBA1001043	C-HEMBA1001043	10546	10547
HEMBA1001052	C-HEMBA1001052	10548	10549
HEMBA1001080	C-HEMBA1001080	10550	
HEMBA1001085	C-HEMBA1001085	10551	10552
HEMBA1001088	C-HEMBA1001088	10553	10554
HEMBA1001109	C-HEMBA1001109	10555	
HEMBA1001122	C-HEMBA1001122	10556	
HEMBA1001133	C-HEMBA1001133	10557	
HEMBA1001137	C-HEMBA1001137	10558	10559
HEMBA1001140	C-HEMBA1001140	10560	10561
HEMBA1001174	C-HEMBA1001174	10562	10563
HEMBA1001197	C-HEMBA1001197	10564	10565
HEMBA1001235	C-HEMBA1001235	10566	
HEMBA1001257	C-HEMBA1001257	10567	10568
HEMBA1001281	C-HEMBA1001281	10569	10570
HEMBA1001286	C-HEMBA1001286	10571	10572
HEMBA1001303	C-HEMBA1001303	10573	
HEMBA1001310	C-HEMBA1001310	10574	10575
HEMBA1001326	C-HEMBA1001326	10576	10577
HEMBA1001351	C-HEMBA1001351	10578	10579
HEMBA1001387	C-HEMBA1001387	10580	10581
HEMBA1001388	C-HEMBA1001388	10582	10583
HEMBA1001398	C-HEMBA1001398	10584	10585
HEMBA1001405	C-HEMBA1001405	10586	10587
HEMBA1001407	C-HEMBA1001407	10588	10589
HEMBA1001413	C-HEMBA1001413	10590	10591
HEMBA1001415	C-HEMBA1001415	10592	10593
HEMBA1001446	C-HEMBA1001446	10594	10595
HEMBA1001450	C-HEMBA1001450	10596	
HEMBA1001455	C-HEMBA1001455	10597	10598
HEMBA1001510	C-HEMBA1001510	10599	10600
HEMBA1001526	C-HEMBA1001526	10601	10602
HEMBA1001533	C-HEMBA1001533	10603	
HEMBA1001579	C-HEMBA1001579	10604	10605
HEMBA1001581	C-HEMBA1001581	10606	
HEMBA1001595	C-HEMBA1001595	10607	10608
HEMBA1001635	C-HEMBA1001635	10609	10610
HEMBA1001661	C-HEMBA1001661	10611	10612
HEMBA1001702	C-HEMBA1001702	10613	
HEMBA1001714	C-HEMBA1001714	10614	
HEMBA1001731	C-HEMBA1001731	10615	
HEMBA1001744	C-HEMBA1001744	10616	10617
HEMBA1001809	C-HEMBA1001809	10618	
HEMBA1001815	C-HEMBA1001815	10619	

(921)

特開 2002-191363

1839

1840

HEMBA1001819	C-HEMBA1001819	10620	10621
HEMBA1001847	C-HEMBA1001847	10622	10623
HEMBA1001864	C-HEMBA1001864	10624	
HEMBA1001869	C-HEMBA1001869	10625	10626
HEMBA1001896	C-HEMBA1001896	10627	10628
HEMBA1001987	C-HEMBA1001987	10629	
HEMBA1002018	C-HEMBA1002018	10630	10631
HEMBA1002049	C-HEMBA1002049	10632	
HEMBA1002084	C-HEMBA1002084	10633	10634
HEMBA1002125	C-HEMBA1002125	10635	10636
HEMBA1002161	C-HEMBA1002161	10637	10638
HEMBA1002177	C-HEMBA1002177	10639	10640
HEMBA1002191	C-HEMBA1002191	10641	
HEMBA1002199	C-HEMBA1002199	10642	10643
HEMBA1002212	C-HEMBA1002212	10644	10645
HEMBA1002237	C-HEMBA1002237	10646	
HEMBA1002265	C-HEMBA1002265	10647	
HEMBA1002267	C-HEMBA1002267	10648	
HEMBA1002349	C-HEMBA1002349	10649	
HEMBA1002363	C-HEMBA1002363	10650	10651
HEMBA1002419	C-HEMBA1002419	10652	10653
HEMBA1002430	C-HEMBA1002430	10654	
HEMBA1002439	C-HEMBA1002439	10655	
HEMBA1002458	C-HEMBA1002458	10656	10657
HEMBA1002460	C-HEMBA1002460	10658	
HEMBA1002462	C-HEMBA1002462	10659	10660
HEMBA1002469	C-HEMBA1002469	10661	10662
HEMBA1002475	C-HEMBA1002475	10663	10664
HEMBA1002477	C-HEMBA1002477	10665	10666
HEMBA1002495	C-HEMBA1002495	10667	10668
HEMBA1002515	C-HEMBA1002515	10669	
HEMBA1002542	C-HEMBA1002542	10670	
HEMBA1002569	C-HEMBA1002569	10671	10672
HEMBA1002583	C-HEMBA1002583	10673	
HEMBA1002609	C-HEMBA1002609	10674	10675
HEMBA1002624	C-HEMBA1002624	10676	10677
HEMBA1002688	C-HEMBA1002688	10678	10679
HEMBA1002696	C-HEMBA1002696	10680	10681
HEMBA1002750	C-HEMBA1002750	10682	
HEMBA1002768	C-HEMBA1002768	10683	10684
HEMBA1002770	C-HEMBA1002770	10685	10686
HEMBA1002777	C-HEMBA1002777	10687	10688
HEMBA1002794	C-HEMBA1002794	10689	10690
HEMBA1002810	C-HEMBA1002810	10691	10692
HEMBA1002818	C-HEMBA1002818	10693	10694
HEMBA1002850	C-HEMBA1002850	10695	
HEMBA1002863	C-HEMBA1002863	10696	10697
HEMBA1002876	C-HEMBA1002876	10698	10699
HEMBA1002935	C-HEMBA1002935	10700	10701
HEMBA1002937	C-HEMBA1002937	10702	10703

(922)

特開 2002-191363

1841

1842

HEMBA1002939	C-HEMBA1002939	10704	10705
HEMBA1002951	C-HEMBA1002951	10706	10707
HEMBA1002954	C-HEMBA1002954	10708	10709
HEMBA1002971	C-HEMBA1002971	10710	
HEMBA1002973	C-HEMBA1002973	10711	10712
HEMBA1002997	C-HEMBA1002997	10713	10714
HEMBA1003033	C-HEMBA1003033	10715	10716
HEMBA1003035	C-HEMBA1003035	10717	
HEMBA1003041	C-HEMBA1003041	10718	10719
HEMBA1003046	C-HEMBA1003046	10720	10721
HEMBA1003067	C-HEMBA1003067	10722	
HEMBA1003096	C-HEMBA1003096	10723	10724
HEMBA1003117	C-HEMBA1003117	10725	10726
HEMBA1003129	C-HEMBA1003129	10727	
HEMBA1003136	C-HEMBA1003136	10728	10729
HEMBA1003148	C-HEMBA1003148	10730	10731
HEMBA1003175	C-HEMBA1003175	10732	10733
HEMBA1003179	C-HEMBA1003179	10734	10735
HEMBA1003199	C-HEMBA1003199	10736	10737
HEMBA1003222	C-HEMBA1003222	10738	10739
HEMBA1003235	C-HEMBA1003235	10740	10741
HEMBA1003250	C-HEMBA1003250	10742	10743
HEMBA1003257	C-HEMBA1003257	10744	10745
HEMBA1003281	C-HEMBA1003281	10746	10747
HEMBA1003286	C-HEMBA1003286	10748	10749
HEMBA1003291	C-HEMBA1003291	10750	10751
HEMBA1003322	C-HEMBA1003322	10752	
HEMBA1003327	C-HEMBA1003327	10753	
HEMBA1003369	C-HEMBA1003369	10754	10755
HEMBA1003370	C-HEMBA1003370	10756	
HEMBA1003380	C-HEMBA1003380	10757	
HEMBA1003395	C-HEMBA1003395	10758	10759
HEMBA1003402	C-HEMBA1003402	10760	
HEMBA1003408	C-HEMBA1003408	10761	10762
HEMBA1003417	C-HEMBA1003417	10763	10764
HEMBA1003418	C-HEMBA1003418	10765	10766
HEMBA1003433	C-HEMBA1003433	10767	10768
HEMBA1003447	C-HEMBA1003447	10769	10770
HEMBA1003461	C-HEMBA1003461	10771	10772
HEMBA1003463	C-HEMBA1003463	10773	
HEMBA1003528	C-HEMBA1003528	10774	10775
HEMBA1003545	C-HEMBA1003545	10776	10777
HEMBA1003555	C-HEMBA1003555	10778	10779
HEMBA1003560	C-HEMBA1003560	10780	
HEMBA1003568	C-HEMBA1003568	10781	10782
HEMBA1003569	C-HEMBA1003569	10783	10784
HEMBA1003581	C-HEMBA1003581	10785	
HEMBA1003591	C-HEMBA1003591	10786	10787
HEMBA1003615	C-HEMBA1003615	10788	10789
HEMBA1003617	C-HEMBA1003617	10790	10791

(923)

特開 2002-191363

1843

1844

HEMBA1003621	C-HEMBA1003621	10792	
HEMBA1003662	C-HEMBA1003662	10793	10794
HEMBA1003690	C-HEMBA1003690	10795	10796
HEMBA1003711	C-HEMBA1003711	10797	10798
HEMBA1003807	C-HEMBA1003807	10799	
HEMBA1003864	C-HEMBA1003864	10800	10801
HEMBA1003953	C-HEMBA1003953	10802	10803
HEMBA1003959	C-HEMBA1003959	10804	
HEMBA1003989	C-HEMBA1003989	10805	10806
HEMBA1004074	C-HEMBA1004074	10807	
HEMBA1004097	C-HEMBA1004097	10808	10809
HEMBA1004146	C-HEMBA1004146	10810	10811
HEMBA1004199	C-HEMBA1004199	10812	10813
HEMBA1004207	C-HEMBA1004207	10814	
HEMBA1004227	C-HEMBA1004227	10815	10816
HEMBA1004246	C-HEMBA1004246	10817	
HEMBA1004276	C-HEMBA1004276	10818	10819
HEMBA1004289	C-HEMBA1004289	10820	10821
HEMBA1004509	C-HEMBA1004509	10822	10823
HEMBA1004534	C-HEMBA1004534	10824	10825
HEMBA1004596	C-HEMBA1004596	10826	10827
HEMBA1004693	C-HEMBA1004693	10828	10829
HEMBA1004736	C-HEMBA1004736	10830	
HEMBA1004753	C-HEMBA1004753	10831	
HEMBA1004756	C-HEMBA1004756	10832	10833
HEMBA1004758	C-HEMBA1004758	10834	10835
HEMBA1004763	C-HEMBA1004763	10836	10837
HEMBA1004768	C-HEMBA1004768	10838	10839
HEMBA1004771	C-HEMBA1004771	10840	
HEMBA1004776	C-HEMBA1004776	10841	
HEMBA1004795	C-HEMBA1004795	10842	10843
HEMBA1004806	C-HEMBA1004806	10844	
HEMBA1004847	C-HEMBA1004847	10845	10846
HEMBA1004850	C-HEMBA1004850	10847	10848
HEMBA1004863	C-HEMBA1004863	10849	
HEMBA1004923	C-HEMBA1004923	10850	10851
HEMBA1004929	C-HEMBA1004929	10852	10853
HEMBA1004930	C-HEMBA1004930	10854	10855
HEMBA1004933	C-HEMBA1004933	10856	10857
HEMBA1004954	C-HEMBA1004954	10858	
HEMBA1004972	C-HEMBA1004972	10859	10860
HEMBA1005475	C-HEMBA1005475	10861	
HEMBA1005581	C-HEMBA1005581	10862	10863
HEMBA1006248	C-HEMBA1006248	10864	10865
HEMBA1006310	C-HEMBA1006310	10866	10867
HEMBA1006344	C-HEMBA1006344	10868	10869
HEMBA1006377	C-HEMBA1006377	10870	10871
HEMBA1006467	C-HEMBA1006467	10872	
HEMBA1006474	C-HEMBA1006474	10873	10874
HEMBA1006530	C-HEMBA1006530	10875	

(924)

特開 2002-191363

1845

1846

HEMBA1006737	C-HEMBA1006737	10876	10877
HEMBA1006795	C-HEMBA1006795	10878	
HEMBA1006877	C-HEMBA1006877	10879	10880
HEMBA1006936	C-HEMBA1006936	10881	10882
HEMBA1007018	C-HEMBA1007018	10883	10884
HEMBA1007342	C-HEMBA1007342	10885	
HEMBA1000008	C-HEMBA1000008	10886	10887
HEMBA1000018	C-HEMBA1000018	10888	
HEMBA1000024	C-HEMBA1000024	10889	
HEMBA1000025	C-HEMBA1000025	10890	
HEMBA1000036	C-HEMBA1000036	10891	10892
HEMBA1000037	C-HEMBA1000037	10893	10894
HEMBA1000083	C-HEMBA1000083	10895	10896
HEMBA1000103	C-HEMBA1000103	10897	
HEMBA1000119	C-HEMBA1000119	10898	10899
HEMBA1000136	C-HEMBA1000136	10900	
HEMBA1000215	C-HEMBA1000215	10901	
HEMBA1000226	C-HEMBA1000226	10902	10903
HEMBA1000244	C-HEMBA1000244	10904	10905
HEMBA1000266	C-HEMBA1000266	10906	10907
HEMBA1000338	C-HEMBA1000338	10908	
HEMBA1000339	C-HEMBA1000339	10909	
HEMBA1000391	C-HEMBA1000391	10910	10911
HEMBA1000438	C-HEMBA1000438	10912	
HEMBA1000449	C-HEMBA1000449	10913	
HEMBA1000589	C-HEMBA1000589	10914	
HEMBA1000591	C-HEMBA1000591	10915	
HEMBA1000623	C-HEMBA1000623	10916	10917
HEMBA1000630	C-HEMBA1000630	10918	10919
HEMBA1000631	C-HEMBA1000631	10920	10921
HEMBA1000632	C-HEMBA1000632	10922	10923
HEMBA1000671	C-HEMBA1000671	10924	
HEMBA1000673	C-HEMBA1000673	10925	10926
HEMBA1000705	C-HEMBA1000705	10927	
HEMBA1000706	C-HEMBA1000706	10928	
HEMBA1000725	C-HEMBA1000725	10929	10930
HEMBA1000763	C-HEMBA1000763	10931	10932
HEMBA1000781	C-HEMBA1000781	10933	10934
HEMBA1000789	C-HEMBA1000789	10935	10936
HEMBA1000807	C-HEMBA1000807	10937	10938
HEMBA1000810	C-HEMBA1000810	10939	
HEMBA1000848	C-HEMBA1000848	10940	10941
HEMBA1000852	C-HEMBA1000852	10942	
HEMBA1000870	C-HEMBA1000870	10943	
HEMBA1000887	C-HEMBA1000887	10944	
HEMBA1000908	C-HEMBA1000908	10945	
HEMBA1000927	C-HEMBA1000927	10946	10947
HEMBA1000947	C-HEMBA1000947	10948	10949
HEMBA1000973	C-HEMBA1000973	10950	10951
HEMBA1000975	C-HEMBA1000975	10952	10953

(925)

特開 2002-191363

1847

1848

HEMBB1000985	C-HEMBB1000985	10954	10955
HEMBB1000991	C-HEMBB1000991	10956	
HEMBB1001011	C-HEMBB1001011	10957	
HEMBB1001014	C-HEMBB1001014	10958	
HEMBB1001024	C-HEMBB1001024	10959	
HEMBB1001056	C-HEMBB1001056	10960	10961
HEMBB1001058	C-HEMBB1001058	10962	10963
HEMBB1001068	C-HEMBB1001068	10964	10965
HEMBB1001096	C-HEMBB1001096	10966	
HEMBB1001105	C-HEMBB1001105	10967	
HEMBB1001117	C-HEMBB1001117	10968	
HEMBB1001126	C-HEMBB1001126	10969	10970
HEMBB1001137	C-HEMBB1001137	10971	10972
HEMBB1001151	C-HEMBB1001151	10973	10974
HEMBB1001153	C-HEMBB1001153	10975	10976
HEMBB1001169	C-HEMBB1001169	10977	
HEMBB1001175	C-HEMBB1001175	10978	10979
HEMBB1001182	C-HEMBB1001182	10980	10981
HEMBB1001199	C-HEMBB1001199	10982	
HEMBB1001210	C-HEMBB1001210	10983	
HEMBB1001242	C-HEMBB1001242	10984	10985
HEMBB1001288	C-HEMBB1001288	10986	10987
HEMBB1001289	C-HEMBB1001289	10988	10989
HEMBB1001294	C-HEMBB1001294	10990	10991
HEMBB1001314	C-HEMBB1001314	10992	10993
HEMBB1001331	C-HEMBB1001331	10994	10995
HEMBB1001339	C-HEMBB1001339	10996	10997
HEMBB1001346	C-HEMBB1001346	10998	10999
HEMBB1001369	C-HEMBB1001369	11000	
HEMBB1001384	C-HEMBB1001384	11001	11002
HEMBB1001387	C-HEMBB1001387	11003	11004
MAMMA1002317	C-MAMMA1002317	11005	
MAMMA1002319	C-MAMMA1002319	11006	
MAMMA1002385	C-MAMMA1002385	11007	11008
NT2RM1000080	C-NT2RM1000080	11009	11010
NT2RM1000242	C-NT2RM1000242	11011	
NT2RM1000257	C-NT2RM1000257	11012	11013
NT2RM1000280	C-NT2RM1000280	11014	11015
NT2RM1000669	C-NT2RM1000669	11016	11017
NT2RM1000781	C-NT2RM1000781	11018	
NT2RM1000867	C-NT2RM1000867	11019	11020
NT2RM1001008	C-NT2RM1001008	11021	11022
NT2RM1001044	C-NT2RM1001044	11023	11024
NT2RM1001074	C-NT2RM1001074	11025	11026
NT2RM1001115	C-NT2RM1001115	11027	11028
NT2RM2000006	C-NT2RM2000006	11029	
NT2RM2000013	C-NT2RM2000013	11030	11031
NT2RM2000030	C-NT2RM2000030	11032	11033
NT2RM2000032	C-NT2RM2000032	11034	
NT2RM2000042	C-NT2RM2000042	11035	

(926)

特開 2002-191363

1849

1850

NT2RM2000092	C-NT2RM2000092	11036	11037
NT2RM2000093	C-NT2RM2000093	11038	11039
NT2RM2000101	C-NT2RM2000101	11040	11041
NT2RM2000191	C-NT2RM2000191	11042	11043
NT2RM2000192	C-NT2RM2000192	11044	11045
NT2RM2000239	C-NT2RM2000239	11046	11047
NT2RM2000250	C-NT2RM2000250	11048	11049
NT2RM2000259	C-NT2RM2000259	11050	11051
NT2RM2000260	C-NT2RM2000260	11052	11053
NT2RM2000287	C-NT2RM2000287	11054	11055
NT2RM2000322	C-NT2RM2000322	11056	11057
NT2RM2000359	C-NT2RM2000359	11058	11059
NT2RM2000363	C-NT2RM2000363	11060	11061
NT2RM2000368	C-NT2RM2000368	11062	11063
NT2RM2000371	C-NT2RM2000371	11064	11065
NT2RM2000374	C-NT2RM2000374	11066	11067
NT2RM2000395	C-NT2RM2000395	11068	11069
NT2RM2000402	C-NT2RM2000402	11070	11071
NT2RM2000407	C-NT2RM2000407	11072	11073
NT2RM2000422	C-NT2RM2000422	11074	11075
NT2RM2000452	C-NT2RM2000452	11076	11077
NT2RM2000469	C-NT2RM2000469	11078	11079
NT2RM2000490	C-NT2RM2000490	11080	11081
NT2RM2000502	C-NT2RM2000502	11082	11083
NT2RM2000504	C-NT2RM2000504	11084	11085
NT2RM2000522	C-NT2RM2000522	11086	11087
NT2RM2000540	C-NT2RM2000540	11088	11089
NT2RM2000567	C-NT2RM2000567	11090	11091
NT2RM2000569	C-NT2RM2000569	11092	
NT2RM2000577	C-NT2RM2000577	11093	11094
NT2RM2000581	C-NT2RM2000581	11095	11096
NT2RM2000588	C-NT2RM2000588	11097	11098
NT2RM2000594	C-NT2RM2000594	11099	11100
NT2RM2000599	C-NT2RM2000599	11101	11102
NT2RM2000624	C-NT2RM2000624	11103	11104
NT2RM2000635	C-NT2RM2000635	11105	11106
NT2RM2000636	C-NT2RM2000636	11107	
NT2RM2000639	C-NT2RM2000639	11108	11109
NT2RM2000649	C-NT2RM2000649	11110	11111
NT2RM2000669	C-NT2RM2000669	11112	11113
NT2RM2000691	C-NT2RM2000691	11114	11115
NT2RM2000714	C-NT2RM2000714	11116	11117
NT2RM2000718	C-NT2RM2000718	11118	11119
NT2RM2000740	C-NT2RM2000740	11120	11121
NT2RM2000795	C-NT2RM2000795	11122	
NT2RM2000821	C-NT2RM2000821	11123	11124
NT2RM2000837	C-NT2RM2000837	11125	11126
NT2RM2000951	C-NT2RM2000951	11127	11128
NT2RM2000952	C-NT2RM2000952	11129	11130
NT2RM2000984	C-NT2RM2000984	11131	11132

1851

1852

NT2RM2001004	C-NT2RM2001004	11133	11134
NT2RM2001035	C-NT2RM2001035	11135	11136
NT2RM2001065	C-NT2RM2001065	11137	11138
NT2RM2001100	C-NT2RM2001100	11139	11140
NT2RM2001131	C-NT2RM2001131	11141	11142
NT2RM2001141	C-NT2RM2001141	11143	11144
NT2RM2001152	C-NT2RM2001152	11145	11146
NT2RM2001177	C-NT2RM2001177	11147	
NT2RM2001194	C-NT2RM2001194	11148	11149
NT2RM2001196	C-NT2RM2001196	11150	11151
NT2RM2001201	C-NT2RM2001201	11152	11153
NT2RM2001221	C-NT2RM2001221	11154	11155
NT2RM2001238	C-NT2RM2001238	11156	11157
NT2RM2001243	C-NT2RM2001243	11158	11159
NT2RM2001247	C-NT2RM2001247	11160	11161
NT2RM2001256	C-NT2RM2001256	11162	11163
NT2RM2001291	C-NT2RM2001291	11164	
NT2RM2001306	C-NT2RM2001306	11165	11166
NT2RM2001312	C-NT2RM2001312	11167	
NT2RM2001319	C-NT2RM2001319	11168	11169
NT2RM2001324	C-NT2RM2001324	11170	11171
NT2RM2001345	C-NT2RM2001345	11172	11173
NT2RM2001370	C-NT2RM2001370	11174	11175
NT2RM2001393	C-NT2RM2001393	11176	11177
NT2RM2001420	C-NT2RM2001420	11178	
NT2RM2001424	C-NT2RM2001424	11179	11180
NT2RM2001499	C-NT2RM2001499	11181	11182
NT2RM2001504	C-NT2RM2001504	11183	11184
NT2RM2001524	C-NT2RM2001524	11185	11186
NT2RM2001544	C-NT2RM2001544	11187	11188
NT2RM2001547	C-NT2RM2001547	11189	11190
NT2RM2001575	C-NT2RM2001575	11191	11192
NT2RM2001582	C-NT2RM2001582	11193	11194
NT2RM2001886	C-NT2RM2001886	11195	11196
NT2RM2001896	C-NT2RM2001896	11197	11198
NT2RM2001903	C-NT2RM2001903	11199	11200
NT2RM2001930	C-NT2RM2001930	11201	11202
NT2RM2001935	C-NT2RM2001935	11203	11204
NT2RM2001936	C-NT2RM2001936	11205	11206
NT2RM2001950	C-NT2RM2001950	11207	11208
NT2RM2001982	C-NT2RM2001982	11209	11210
NT2RM2001989	C-NT2RM2001989	11211	11212
NT2RM2001997	C-NT2RM2001997	11213	11214
NT2RM2001998	C-NT2RM2001998	11215	11216
NT2RM2002004	C-NT2RM2002004	11217	11218
NT2RM2002014	C-NT2RM2002014	11219	11220
NT2RM2002030	C-NT2RM2002030	11221	11222
NT2RM2002049	C-NT2RM2002049	11223	11224
NT2RM2002055	C-NT2RM2002055	11225	11226
NT2RM2002088	C-NT2RM2002088	11227	11228

1853

1854

NT2RM2002091	C-NT2RM2002091	11229	11230
NT2RM2002100	C-NT2RM2002100	11231	11232
NT2RM2002109	C-NT2RM2002109	11233	11234
NT2RM2002128	C-NT2RM2002128	11235	11236
NT2RM2002142	C-NT2RM2002142	11237	11238
NT2RM2002178	C-NT2RM2002178	11239	11240
NT2RM4000024	C-NT2RM4000024	11241	11242
NT2RM4000061	C-NT2RM4000061	11243	
NT2RM4000104	C-NT2RM4000104	11244	11245
NT2RM4000139	C-NT2RM4000139	11246	11247
NT2RM4000169	C-NT2RM4000169	11248	11249
NT2RM4000191	C-NT2RM4000191	11250	11251
NT2RM4000197	C-NT2RM4000197	11252	
NT2RM4000210	C-NT2RM4000210	11253	11254
NT2RM4000229	C-NT2RM4000229	11255	11256
NT2RM4000290	C-NT2RM4000290	11257	11258
NT2RM4000344	C-NT2RM4000344	11259	11260
NT2RM4000349	C-NT2RM4000349	11261	11262
NT2RM4000354	C-NT2RM4000354	11263	11264
NT2RM4000386	C-NT2RM4000386	11265	11266
NT2RM4000395	C-NT2RM4000395	11267	11268
NT2RM4000421	C-NT2RM4000421	11269	11270
NT2RM4000457	C-NT2RM4000457	11271	11272
NT2RM4000471	C-NT2RM4000471	11273	11274
NT2RM4000486	C-NT2RM4000486	11275	11276
NT2RM4000496	C-NT2RM4000496	11277	11278
NT2RM4000511	C-NT2RM4000511	11279	11280
NT2RM4000515	C-NT2RM4000515	11281	11282
NT2RM4000520	C-NT2RM4000520	11283	11284
NT2RM4000585	C-NT2RM4000585	11285	11286
NT2RM4000595	C-NT2RM4000595	11287	11288
NT2RP1000018	C-NT2RP1000018	11289	11290
NT2RP1000035	C-NT2RP1000035	11291	11292
NT2RP1000040	C-NT2RP1000040	11293	11294
NT2RP1000063	C-NT2RP1000063	11295	11296
NT2RP1000086	C-NT2RP1000086	11297	11298
NT2RP1000101	C-NT2RP1000101	11299	11300
NT2RP1000111	C-NT2RP1000111	11301	11302
NT2RP1000112	C-NT2RP1000112	11303	
NT2RP1000124	C-NT2RP1000124	11304	11305
NT2RP1000130	C-NT2RP1000130	11306	11307
NT2RP1000163	C-NT2RP1000163	11308	11309
NT2RP1000170	C-NT2RP1000170	11310	11311
NT2RP1000191	C-NT2RP1000191	11312	11313
NT2RP1000202	C-NT2RP1000202	11314	11315
NT2RP1000243	C-NT2RP1000243	11316	11317
NT2RP1000259	C-NT2RP1000259	11318	
NT2RP1000272	C-NT2RP1000272	11319	11320
NT2RP1000326	C-NT2RP1000326	11321	11322
NT2RP1000333	C-NT2RP1000333	11323	11324

1855

1856

NT2RP1000348	C-NT2RP1000348	11325	11326
NT2RP1000357	C-NT2RP1000357	11327	11328
NT2RP1000376	C-NT2RP1000376	11329	11330
NT2RP1000413	C-NT2RP1000413	11331	11332
NT2RP1000416	C-NT2RP1000416	11333	11334
NT2RP1000439	C-NT2RP1000439	11335	11336
NT2RP1000443	C-NT2RP1000443	11337	11338
NT2RP1000470	C-NT2RP1000470	11339	11340
NT2RP1000478	C-NT2RP1000478	11341	11342
NT2RP1000481	C-NT2RP1000481	11343	11344
NT2RP1000493	C-NT2RP1000493	11345	11346
NT2RP1000547	C-NT2RP1000547	11347	11348
NT2RP1000574	C-NT2RP1000574	11349	11350
NT2RP1000581	C-NT2RP1000581	11351	
NT2RP1000630	C-NT2RP1000630	11352	11353
NT2RP1000688	C-NT2RP1000688	11354	11355
NT2RP1000695	C-NT2RP1000695	11356	11357
NT2RP1000733	C-NT2RP1000733	11358	11359
NT2RP1000738	C-NT2RP1000738	11360	11361
NT2RP1000782	C-NT2RP1000782	11362	11363
NT2RP1000825	C-NT2RP1000825	11364	11365
NT2RP1000833	C-NT2RP1000833	11366	11367
NT2RP1000846	C-NT2RP1000846	11368	
NT2RP1000851	C-NT2RP1000851	11369	
NT2RP1000856	C-NT2RP1000856	11370	11371
NT2RP1000915	C-NT2RP1000915	11372	11373
NT2RP1000947	C-NT2RP1000947	11374	11375
NT2RP1000954	C-NT2RP1000954	11376	11377
NT2RP1000958	C-NT2RP1000958	11378	11379
NT2RP1000959	C-NT2RP1000959	11380	11381
NT2RP1000966	C-NT2RP1000966	11382	11383
NT2RP1000980	C-NT2RP1000980	11384	11385
NT2RP1000988	C-NT2RP1000988	11386	11387
NT2RP1001011	C-NT2RP1001011	11388	11389
NT2RP1001014	C-NT2RP1001014	11390	11391
NT2RP1001395	C-NT2RP1001395	11392	11393
NT2RP1001410	C-NT2RP1001410	11394	11395
NT2RP1001424	C-NT2RP1001424	11396	
NT2RP1001449	C-NT2RP1001449	11397	11398
NT2RP1001457	C-NT2RP1001457	11399	11400
NT2RP1001466	C-NT2RP1001466	11401	11402
NT2RP1001475	C-NT2RP1001475	11403	11404
NT2RP1001482	C-NT2RP1001482	11405	11406
NT2RP1001494	C-NT2RP1001494	11407	11408
NT2RP1001543	C-NT2RP1001543	11409	11410
NT2RP1001546	C-NT2RP1001546	11411	11412
NT2RP1001569	C-NT2RP1001569	11413	11414
NT2RP1001616	C-NT2RP1001616	11415	
NT2RP1001665	C-NT2RP1001665	11416	11417
NT2RP2000006	C-NT2RP2000006	11418	

(930)

特開 2002-191363

1857

1858

NT2RP2000007	C-NT2RP2000007	11419	11420
NT2RP2000008	C-NT2RP2000008	11421	11422
NT2RP2000032	C-NT2RP2000032	11423	11424
NT2RP2000045	C-NT2RP2000045	11425	11426
NT2RP2000054	C-NT2RP2000054	11427	11428
NT2RP2000056	C-NT2RP2000056	11429	
NT2RP2000067	C-NT2RP2000067	11430	11431
NT2RP2000070	C-NT2RP2000070	11432	11433
NT2RP2000079	C-NT2RP2000079	11434	11435
NT2RP2000088	C-NT2RP2000088	11436	
NT2RP2000091	C-NT2RP2000091	11437	11438
NT2RP2000097	C-NT2RP2000097	11439	11440
NT2RP2000114	C-NT2RP2000114	11441	11442
NT2RP2000120	C-NT2RP2000120	11443	11444
NT2RP2000126	C-NT2RP2000126	11445	11446
NT2RP2000133	C-NT2RP2000133	11447	11448
NT2RP2000147	C-NT2RP2000147	11449	11450
NT2RP2000153	C-NT2RP2000153	11451	11452
NT2RP2000157	C-NT2RP2000157	11453	11454
NT2RP2000161	C-NT2RP2000161	11455	11456
NT2RP2000173	C-NT2RP2000173	11457	11458
NT2RP2000175	C-NT2RP2000175	11459	11460
NT2RP2000195	C-NT2RP2000195	11461	11462
NT2RP2000205	C-NT2RP2000205	11463	11464
NT2RP2000208	C-NT2RP2000208	11465	11466
NT2RP2000224	C-NT2RP2000224	11467	11468
NT2RP2000232	C-NT2RP2000232	11469	
NT2RP2000233	C-NT2RP2000233	11470	11471
NT2RP2000239	C-NT2RP2000239	11472	11473
NT2RP2000248	C-NT2RP2000248	11474	11475
NT2RP2000270	C-NT2RP2000270	11476	11477
NT2RP2000274	C-NT2RP2000274	11478	11479
NT2RP2000283	C-NT2RP2000283	11480	11481
NT2RP2000288	C-NT2RP2000288	11482	11483
NT2RP2000297	C-NT2RP2000297	11484	
NT2RP2000298	C-NT2RP2000298	11485	11486
NT2RP2000310	C-NT2RP2000310	11487	11488
NT2RP2000328	C-NT2RP2000328	11489	11490
NT2RP2000329	C-NT2RP2000329	11491	11492
NT2RP2000346	C-NT2RP2000346	11493	11494
NT2RP2000369	C-NT2RP2000369	11495	
NT2RP2000412	C-NT2RP2000412	11496	11497
NT2RP2000414	C-NT2RP2000414	11498	11499
NT2RP2000422	C-NT2RP2000422	11500	11501
NT2RP2000438	C-NT2RP2000438	11502	11503
NT2RP2000448	C-NT2RP2000448	11504	11505
NT2RP2000503	C-NT2RP2000503	11506	
NT2RP2000510	C-NT2RP2000510	11507	11508
NT2RP2000516	C-NT2RP2000516	11509	11510
NT2RP2000603	C-NT2RP2000603	11511	11512

(931)

特開 2002-191363

1859

1860

NT2RP2000617	C-NT2RP2000617	11513	11514
NT2RP2000634	C-NT2RP2000634	11515	11516
NT2RP2000656	C-NT2RP2000656	11517	11518
NT2RP2000658	C-NT2RP2000658	11519	11520
NT2RP2000668	C-NT2RP2000668	11521	11522
NT2RP2000704	C-NT2RP2000704	11523	
NT2RP2000710	C-NT2RP2000710	11524	11525
NT2RP2000764	C-NT2RP2000764	11526	11527
NT2RP2000809	C-NT2RP2000809	11528	11529
NT2RP2000812	C-NT2RP2000812	11530	11531
NT2RP2000814	C-NT2RP2000814	11532	11533
NT2RP2000816	C-NT2RP2000816	11534	11535
NT2RP2000819	C-NT2RP2000819	11536	11537
NT2RP2000841	C-NT2RP2000841	11538	11539
NT2RP2000845	C-NT2RP2000845	11540	
NT2RP2000863	C-NT2RP2000863	11541	11542
NT2RP2000880	C-NT2RP2000880	11543	11544
NT2RP2000892	C-NT2RP2000892	11545	11546
NT2RP2000931	C-NT2RP2000931	11547	11548
NT2RP2000932	C-NT2RP2000932	11549	11550
NT2RP2000938	C-NT2RP2000938	11551	11552
NT2RP2000943	C-NT2RP2000943	11553	11554
NT2RP2000965	C-NT2RP2000965	11555	11556
NT2RP2000985	C-NT2RP2000985	11557	11558
NT2RP2001036	C-NT2RP2001036	11559	
NT2RP2001044	C-NT2RP2001044	11560	
NT2RP2001056	C-NT2RP2001056	11561	
NT2RP2001065	C-NT2RP2001065	11562	11563
NT2RP2001070	C-NT2RP2001070	11564	11565
NT2RP2001081	C-NT2RP2001081	11566	11567
NT2RP2001094	C-NT2RP2001094	11568	
NT2RP2001119	C-NT2RP2001119	11569	11570
NT2RP2001127	C-NT2RP2001127	11571	11572
NT2RP2001218	C-NT2RP2001218	11573	11574
NT2RP2001245	C-NT2RP2001245	11575	11576
NT2RP2001381	C-NT2RP2001381	11577	11578
NT2RP2001397	C-NT2RP2001397	11579	11580
NT2RP2001427	C-NT2RP2001427	11581	
NT2RP2001601	C-NT2RP2001601	11582	11583
NT2RP2001675	C-NT2RP2001675	11584	11585
NT2RP2001721	C-NT2RP2001721	11586	11587
NT2RP2001907	C-NT2RP2001907	11588	11589
NT2RP2001969	C-NT2RP2001969	11590	11591
NT2RP2001976	C-NT2RP2001976	11592	11593
NT2RP2002046	C-NT2RP2002046	11594	11595
NT2RP2002154	C-NT2RP2002154	11596	11597
NT2RP2002208	C-NT2RP2002208	11598	11599
NT2RP2002270	C-NT2RP2002270	11600	11601
NT2RP2002312	C-NT2RP2002312	11602	11603
NT2RP2002325	C-NT2RP2002325	11604	11605

1861

1862

NT2RP2002385	C-NT2RP2002385	11606	11607
NT2RP2002426	C-NT2RP2002426	11608	
NT2RP2002479	C-NT2RP2002479	11609	11610
NT2RP2002537	C-NT2RP2002537	11611	11612
NT2RP2002595	C-NT2RP2002595	11613	11614
NT2RP2002618	C-NT2RP2002618	11615	11616
NT2RP2002621	C-NT2RP2002621	11617	11618
NT2RP2002672	C-NT2RP2002672	11619	
NT2RP2002701	C-NT2RP2002701	11620	11621
NT2RP2002769	C-NT2RP2002769	11622	11623
NT2RP2002862	C-NT2RP2002862	11624	11625
NT2RP2002928	C-NT2RP2002928	11626	11627
NT2RP2002954	C-NT2RP2002954	11628	11629
NT2RP2002959	C-NT2RP2002959	11630	11631
NT2RP2002980	C-NT2RP2002980	11632	11633
NT2RP2002986	C-NT2RP2002986	11634	11635
NT2RP2003108	C-NT2RP2003108	11636	11637
NT2RP2003117	C-NT2RP2003117	11638	
NT2RP2003121	C-NT2RP2003121	11639	11640
NT2RP2003125	C-NT2RP2003125	11641	11642
NT2RP2003177	C-NT2RP2003177	11643	11644
NT2RP2003194	C-NT2RP2003194	11645	11646
NT2RP2003265	C-NT2RP2003265	11647	11648
NT2RP2003295	C-NT2RP2003295	11649	11650
NT2RP2003329	C-NT2RP2003329	11651	11652
NT2RP2003367	C-NT2RP2003367	11653	
NT2RP2003433	C-NT2RP2003433	11654	11655
NT2RP2003446	C-NT2RP2003446	11656	11657
NT2RP2003533	C-NT2RP2003533	11658	
NT2RP2003543	C-NT2RP2003543	11659	11660
NT2RP2003596	C-NT2RP2003596	11661	11662
NT2RP2003629	C-NT2RP2003629	11663	11664
NT2RP2003687	C-NT2RP2003687	11665	
NT2RP2003714	C-NT2RP2003714	11666	11667
NT2RP2003737	C-NT2RP2003737	11668	11669
NT2RP2003793	C-NT2RP2003793	11670	11671
NT2RP2003952	C-NT2RP2003952	11672	11673
NT2RP2003986	C-NT2RP2003986	11674	
NT2RP2004042	C-NT2RP2004042	11675	11676
NT2RP2004316	C-NT2RP2004316	11677	
NT2RP2004389	C-NT2RP2004389	11678	11679
NT2RP2004392	C-NT2RP2004392	11680	11681
NT2RP2004463	C-NT2RP2004463	11682	11683
NT2RP2004602	C-NT2RP2004602	11684	
NT2RP2004614	C-NT2RP2004614	11685	11686
NT2RP2004655	C-NT2RP2004655	11687	11688
NT2RP2004689	C-NT2RP2004689	11689	11690
NT2RP2004791	C-NT2RP2004791	11691	11692
NT2RP2004799	C-NT2RP2004799	11693	11694
NT2RP2004802	C-NT2RP2004802	11695	11696

(933)

特開 2002-191363

1863

1864

NT2RP2004841	C-NT2RP2004841	11697	11698
NT2RP2004936	C-NT2RP2004936	11699	11700
NT2RP2004959	C-NT2RP2004959	11701	11702
NT2RP2004999	C-NT2RP2004999	11703	11704
NT2RP2005000	C-NT2RP2005000	11705	11706
NT2RP2005001	C-NT2RP2005001	11707	11708
NT2RP2005012	C-NT2RP2005012	11709	11710
NT2RP2005037	C-NT2RP2005037	11711	11712
NT2RP2005126	C-NT2RP2005126	11713	11714
NT2RP2005140	C-NT2RP2005140	11715	11716
NT2RP2005147	C-NT2RP2005147	11717	
NT2RP2005159	C-NT2RP2005159	11718	11719
NT2RP2005239	C-NT2RP2005239	11720	11721
NT2RP2005270	C-NT2RP2005270	11722	11723
NT2RP2005276	C-NT2RP2005276	11724	11725
NT2RP2005293	C-NT2RP2005293	11726	11727
NT2RP2005315	C-NT2RP2005315	11728	11729
NT2RP2005358	C-NT2RP2005358	11730	11731
NT2RP2005393	C-NT2RP2005393	11732	11733
NT2RP2005436	C-NT2RP2005436	11734	11735
NT2RP2005441	C-NT2RP2005441	11736	11737
NT2RP2005453	C-NT2RP2005453	11738	
NT2RP2005464	C-NT2RP2005464	11739	
NT2RP2005465	C-NT2RP2005465	11740	11741
NT2RP2005472	C-NT2RP2005472	11742	11743
NT2RP2005495	C-NT2RP2005495	11744	11745
NT2RP2005498	C-NT2RP2005498	11746	11747
NT2RP2005509	C-NT2RP2005509	11748	11749
NT2RP2005520	C-NT2RP2005520	11750	11751
NT2RP2005525	C-NT2RP2005525	11752	11753
NT2RP2005540	C-NT2RP2005540	11754	11755
NT2RP2005549	C-NT2RP2005549	11756	11757
NT2RP2005555	C-NT2RP2005555	11758	
NT2RP2005557	C-NT2RP2005557	11759	11760
NT2RP2005620	C-NT2RP2005620	11761	11762
NT2RP2005622	C-NT2RP2005622	11763	11764
NT2RP2005635	C-NT2RP2005635	11765	11766
NT2RP2005637	C-NT2RP2005637	11767	
NT2RP2005640	C-NT2RP2005640	11768	11769
NT2RP2005654	C-NT2RP2005654	11770	11771
NT2RP2005669	C-NT2RP2005669	11772	11773
NT2RP2005675	C-NT2RP2005675	11774	11775
NT2RP2005683	C-NT2RP2005683	11776	11777
NT2RP2005690	C-NT2RP2005690	11778	11779
NT2RP2005712	C-NT2RP2005712	11780	11781
NT2RP2005723	C-NT2RP2005723	11782	11783
NT2RP2005748	C-NT2RP2005748	11784	11785
NT2RP2005752	C-NT2RP2005752	11786	11787
NT2RP2005753	C-NT2RP2005753	11788	11789
NT2RP2005763	C-NT2RP2005763	11790	11791

(934)

特開 2002-191363

1865

1866

NT2RP2005767	C-NT2RP2005767	11792	11793
NT2RP2005773	C-NT2RP2005773	11794	11795
NT2RP2005775	C-NT2RP2005775	11796	11797
NT2RP2005781	C-NT2RP2005781	11798	11799
NT2RP2005804	C-NT2RP2005804	11800	11801
NT2RP2005835	C-NT2RP2005835	11802	11803
NT2RP2005853	C-NT2RP2005853	11804	11805
NT2RP2005868	C-NT2RP2005868	11806	11807
NT2RP2005886	C-NT2RP2005886	11808	11809
NT2RP2005890	C-NT2RP2005890	11810	
NT2RP2005901	C-NT2RP2005901	11811	11812
NT2RP2005933	C-NT2RP2005933	11813	11814
NT2RP2006038	C-NT2RP2006038	11815	11816
NT2RP2006043	C-NT2RP2006043	11817	11818
NT2RP2006052	C-NT2RP2006052	11819	
NT2RP2006069	C-NT2RP2006069	11820	11821
NT2RP2006071	C-NT2RP2006071	11822	11823
NT2RP2006100	C-NT2RP2006100	11824	11825
NT2RP2006106	C-NT2RP2006106	11826	11827
NT2RP2006141	C-NT2RP2006141	11828	11829
NT2RP2006186	C-NT2RP2006186	11830	11831
NT2RP2006196	C-NT2RP2006196	11832	
NT2RP2006200	C-NT2RP2006200	11833	11834
NT2RP2006219	C-NT2RP2006219	11835	11836
NT2RP2006237	C-NT2RP2006237	11837	11838
NT2RP2006238	C-NT2RP2006238	11839	11840
NT2RP2006275	C-NT2RP2006275	11841	11842
NT2RP2006312	C-NT2RP2006312	11843	11844
NT2RP2006333	C-NT2RP2006333	11845	11846
NT2RP2006365	C-NT2RP2006365	11847	11848
NT2RP2006393	C-NT2RP2006393	11849	
NT2RP2006436	C-NT2RP2006436	11850	
NT2RP2006456	C-NT2RP2006456	11851	11852
NT2RP2006464	C-NT2RP2006464	11853	11854
NT2RP2006467	C-NT2RP2006467	11855	
NT2RP2006472	C-NT2RP2006472	11856	11857
NT2RP2006565	C-NT2RP2006565	11858	
NT2RP2006571	C-NT2RP2006571	11859	11860
NT2RP2006573	C-NT2RP2006573	11861	11862
NT2RP3000031	C-NT2RP3000031	11863	11864
NT2RP3000072	C-NT2RP3000072	11865	11866
NT2RP3000142	C-NT2RP3000142	11867	11868
NT2RP3000220	C-NT2RP3000220	11869	
NT2RP3000251	C-NT2RP3000251	11870	
NT2RP3000252	C-NT2RP3000252	11871	11872
NT2RP3000312	C-NT2RP3000312	11873	11874
NT2RP3000320	C-NT2RP3000320	11875	11876
NT2RP3000333	C-NT2RP3000333	11877	11878
NT2RP3000348	C-NT2RP3000348	11879	
NT2RP3000350	C-NT2RP3000350	11880	11881

(935)

特開 2002-191363

1867

1868

NT2RP3000359	C-NT2RP3000359	11882	11883
NT2RP3000361	C-NT2RP3000361	11884	11885
NT2RP3000366	C-NT2RP3000366	11886	11887
NT2RP3000397	C-NT2RP3000397	11888	11889
NT2RP3000403	C-NT2RP3000403	11890	11891
NT2RP3000484	C-NT2RP3000484	11892	
NT2RP3000527	C-NT2RP3000527	11893	11894
NT2RP3000531	C-NT2RP3000531	11895	11896
NT2RP3000596	C-NT2RP3000596	11897	11898
NT2RP3000599	C-NT2RP3000599	11899	11900
NT2RP3000632	C-NT2RP3000632	11901	11902
NT2RP3000644	C-NT2RP3000644	11903	11904
NT2RP3000661	C-NT2RP3000661	11905	11906
NT2RP3000665	C-NT2RP3000665	11907	11908
NT2RP3000690	C-NT2RP3000690	11909	11910
NT2RP3000759	C-NT2RP3000759	11911	11912
NT2RP3000825	C-NT2RP3000825	11913	11914
NT2RP3000836	C-NT2RP3000836	11915	
NT2RP3000841	C-NT2RP3000841	11916	11917
NT2RP3000850	C-NT2RP3000850	11918	
NT2RP3000852	C-NT2RP3000852	11919	11920
NT2RP3000859	C-NT2RP3000859	11921	11922
NT2RP3000868	C-NT2RP3000868	11923	11924
NT2RP3000869	C-NT2RP3000869	11925	11926
NT2RP3000901	C-NT2RP3000901	11927	
NT2RP3000917	C-NT2RP3000917	11928	11929
NT2RP3000919	C-NT2RP3000919	11930	11931
NT2RP3000980	C-NT2RP3000980	11932	11933
NT2RP3000994	C-NT2RP3000994	11934	11935
NT2RP3001004	C-NT2RP3001004	11936	11937
NT2RP3001081	C-NT2RP3001081	11938	11939
NT2RP3001084	C-NT2RP3001084	11940	11941
NT2RP3001096	C-NT2RP3001096	11942	11943
NT2RP3001107	C-NT2RP3001107	11944	11945
NT2RP3001109	C-NT2RP3001109	11946	11947
NT2RP3001116	C-NT2RP3001116	11948	
NT2RP3001119	C-NT2RP3001119	11949	11950
NT2RP3001133	C-NT2RP3001133	11951	11952
NT2RP3001140	C-NT2RP3001140	11953	11954
NT2RP3001155	C-NT2RP3001155	11955	11956
NT2RP3001176	C-NT2RP3001176	11957	11958
NT2RP3001214	C-NT2RP3001214	11959	11960
NT2RP3001216	C-NT2RP3001216	11961	11962
NT2RP3001221	C-NT2RP3001221	11963	11964
NT2RP3001236	C-NT2RP3001236	11965	11966
NT2RP3001239	C-NT2RP3001239	11967	11968
NT2RP3001260	C-NT2RP3001260	11969	11970
NT2RP3001307	C-NT2RP3001307	11971	11972
NT2RP3001325	C-NT2RP3001325	11973	11974
NT2RP3001384	C-NT2RP3001384	11975	11976

(936)

特開 2002-191363

1869

1870

NT2RP3001392	C-NT2RP3001392	11977	
NT2RP3001396	C-NT2RP3001396	11978	11979
NT2RP3001398	C-NT2RP3001398	11980	11981
NT2RP3001407	C-NT2RP3001407	11982	11983
NT2RP3001420	C-NT2RP3001420	11984	11985
NT2RP3001426	C-NT2RP3001426	11986	11987
NT2RP3001427	C-NT2RP3001427	11988	11989
NT2RP3001457	C-NT2RP3001457	11990	11991
NT2RP3001472	C-NT2RP3001472	11992	11993
NT2RP3001495	C-NT2RP3001495	11994	11995
NT2RP3001497	C-NT2RP3001497	11996	11997
NT2RP3001529	C-NT2RP3001529	11998	11999
NT2RP3001621	C-NT2RP3001621	12000	12001
NT2RP3001629	C-NT2RP3001629	12002	
NT2RP3001642	C-NT2RP3001642	12003	12004
NT2RP3001646	C-NT2RP3001646	12005	12006
NT2RP3001676	C-NT2RP3001676	12007	
NT2RP3001679	C-NT2RP3001679	12008	12009
NT2RP3001799	C-NT2RP3001799	12010	12011
NT2RP3001819	C-NT2RP3001819	12012	12013
NT2RP3001896	C-NT2RP3001896	12014	12015
NT2RP3001915	C-NT2RP3001915	12016	12017
NT2RP3001929	C-NT2RP3001929	12018	
NT2RP3003193	C-NT2RP3003193	12019	
NT2RP3004466	C-NT2RP3004466	12020	12021
NT2RP3004480	C-NT2RP3004480	12022	12023
NT2RP3004539	C-NT2RP3004539	12024	12025
NT2RP3004544	C-NT2RP3004544	12026	12027
NT2RP3004569	C-NT2RP3004569	12028	12029
NT2RP3004572	C-NT2RP3004572	12030	12031
NT2RP3004578	C-NT2RP3004578	12032	12033
NT2RP3004594	C-NT2RP3004594	12034	12035
NT2RP3004617	C-NT2RP3004617	12036	12037
NT2RP3004618	C-NT2RP3004618	12038	12039
NT2RP3004669	C-NT2RP3004669	12040	12041
NT2RP4000008	C-NT2RP4000008	12042	12043
NT2RP4000051	C-NT2RP4000051	12044	12045
NT2RP4000078	C-NT2RP4000078	12046	12047
NT2RP4000109	C-NT2RP4000109	12048	
NT2RP4000111	C-NT2RP4000111	12049	12050
NT2RP4000129	C-NT2RP4000129	12051	12052
NT2RP4000147	C-NT2RP4000147	12053	12054
NT2RP4000150	C-NT2RP4000150	12055	
NT2RP4000151	C-NT2RP4000151	12056	12057
NT2RP4000159	C-NT2RP4000159	12058	12059
NT2RP4000185	C-NT2RP4000185	12060	12061
NT2RP4000210	C-NT2RP4000210	12062	12063
NT2RP4000212	C-NT2RP4000212	12064	12065
NT2RP4000243	C-NT2RP4000243	12066	12067
NT2RP4000246	C-NT2RP4000246	12068	12069

(937)

特開 2002-191363

1871

1872

NT2RP4000259	C-NT2RP4000259	12070	12071
NT2RP4000290	C-NT2RP4000290	12072	12073
NT2RP4000312	C-NT2RP4000312	12074	12075
NT2RP4000323	C-NT2RP4000323	12076	12077
NT2RP4000355	C-NT2RP4000355	12078	12079
NT2RP4000360	C-NT2RP4000360	12080	12081
NT2RP4000367	C-NT2RP4000367	12082	12083
NT2RP4000370	C-NT2RP4000370	12084	12085
NT2RP4000376	C-NT2RP4000376	12086	12087
NT2RP4000381	C-NT2RP4000381	12088	12089
NT2RP4000398	C-NT2RP4000398	12090	12091
NT2RP4000415	C-NT2RP4000415	12092	12093
NT2RP4000417	C-NT2RP4000417	12094	12095
NT2RP4000448	C-NT2RP4000448	12096	
NT2RP4000449	C-NT2RP4000449	12097	12098
NT2RP4000455	C-NT2RP4000455	12099	12100
NT2RP4000457	C-NT2RP4000457	12101	12102
NT2RP4000480	C-NT2RP4000480	12103	12104
NT2RP4000481	C-NT2RP4000481	12105	12106
NT2RP4000498	C-NT2RP4000498	12107	12108
NT2RP4000500	C-NT2RP4000500	12109	12110
NT2RP4000518	C-NT2RP4000518	12111	12112
NT2RP4000524	C-NT2RP4000524	12113	12114
NT2RP4000541	C-NT2RP4000541	12115	12116
NT2RP4000556	C-NT2RP4000556	12117	12118
NT2RP4000560	C-NT2RP4000560	12119	12120
NT2RP4000588	C-NT2RP4000588	12121	12122
NT2RP4000614	C-NT2RP4000614	12123	12124
NT2RP4000638	C-NT2RP4000638	12125	12126
NT2RP4000648	C-NT2RP4000648	12127	12128
NT2RP4000657	C-NT2RP4000657	12129	12130
NT2RP4000704	C-NT2RP4000704	12131	12132
NT2RP4000713	C-NT2RP4000713	12133	12134
NT2RP4000724	C-NT2RP4000724	12135	12136
NT2RP4000728	C-NT2RP4000728	12137	12138
NT2RP4000737	C-NT2RP4000737	12139	12140
NT2RP4000739	C-NT2RP4000739	12141	12142
NT2RP4000781	C-NT2RP4000781	12143	12144
NT2RP4000817	C-NT2RP4000817	12145	12146
NT2RP4000833	C-NT2RP4000833	12147	12148
NT2RP4000837	C-NT2RP4000837	12149	12150
NT2RP4000839	C-NT2RP4000839	12151	12152
NT2RP4000855	C-NT2RP4000855	12153	12154
NT2RP4000865	C-NT2RP4000865	12155	12156
NT2RP4000878	C-NT2RP4000878	12157	12158
NT2RP4000879	C-NT2RP4000879	12159	12160
NT2RP4000925	C-NT2RP4000925	12161	12162
NT2RP4000927	C-NT2RP4000927	12163	12164
NT2RP4000928	C-NT2RP4000928	12165	12166
NT2RP4000929	C-NT2RP4000929	12167	12168

(938)

特開 2002-191363

1873

1874

NT2RP4000955	C-NT2RP4000955	12169	12170
NT2RP4000973	C-NT2RP4000973	12171	12172
NT2RP4000975	C-NT2RP4000975	12173	12174
NT2RP4000979	C-NT2RP4000979	12175	12176
NT2RP4000984	C-NT2RP4000984	12177	12178
NT2RP4000989	C-NT2RP4000989	12179	12180
NT2RP4000997	C-NT2RP4000997	12181	12182
NT2RP4001004	C-NT2RP4001004	12183	12184
NT2RP4001006	C-NT2RP4001006	12185	12186
NT2RP4001010	C-NT2RP4001010	12187	12188
NT2RP4001041	C-NT2RP4001041	12189	12190
NT2RP4001057	C-NT2RP4001057	12191	12192
NT2RP4001064	C-NT2RP4001064	12193	12194
NT2RP4001079	C-NT2RP4001079	12195	12196
NT2RP4001080	C-NT2RP4001080	12197	12198
NT2RP4001086	C-NT2RP4001086	12199	12200
NT2RP4001095	C-NT2RP4001095	12201	12202
NT2RP4001100	C-NT2RP4001100	12203	12204
NT2RP4001117	C-NT2RP4001117	12205	12206
NT2RP4001122	C-NT2RP4001122	12207	12208
NT2RP4001126	C-NT2RP4001126	12209	12210
NT2RP4001138	C-NT2RP4001138	12211	12212
NT2RP4001143	C-NT2RP4001143	12213	12214
NT2RP4001148	C-NT2RP4001148	12215	
NT2RP4001149	C-NT2RP4001149	12216	12217
NT2RP4001150	C-NT2RP4001150	12218	12219
NT2RP4001174	C-NT2RP4001174	12220	12221
NT2RP4001206	C-NT2RP4001206	12222	12223
NT2RP4001207	C-NT2RP4001207	12224	12225
NT2RP4001210	C-NT2RP4001210	12226	12227
NT2RP4001219	C-NT2RP4001219	12228	12229
NT2RP4001228	C-NT2RP4001228	12230	12231
NT2RP4001235	C-NT2RP4001235	12232	12233
NT2RP4001256	C-NT2RP4001256	12234	12235
NT2RP4001260	C-NT2RP4001260	12236	12237
NT2RP4001274	C-NT2RP4001274	12238	12239
NT2RP4001276	C-NT2RP4001276	12240	12241
NT2RP4001313	C-NT2RP4001313	12242	12243
NT2RP4001315	C-NT2RP4001315	12244	12245
NT2RP4001339	C-NT2RP4001339	12246	12247
NT2RP4001343	C-NT2RP4001343	12248	12249
NT2RP4001345	C-NT2RP4001345	12250	12251
NT2RP4001351	C-NT2RP4001351	12252	12253
NT2RP4001353	C-NT2RP4001353	12254	12255
NT2RP4001372	C-NT2RP4001372	12256	12257
NT2RP4001373	C-NT2RP4001373	12258	12259
NT2RP4001375	C-NT2RP4001375	12260	12261
NT2RP4001379	C-NT2RP4001379	12262	12263
NT2RP4001407	C-NT2RP4001407	12264	12265
NT2RP4001414	C-NT2RP4001414	12266	12267

(939)

特開 2002-191363

1875

1876

NT2RP4001433	C-NT2RP4001433	12268	12269
NT2RP4001474	C-NT2RP4001474	12270	12271
NT2RP4001483	C-NT2RP4001483	12272	12273
NT2RP4001498	C-NT2RP4001498	12274	12275
NT2RP4001502	C-NT2RP4001502	12276	12277
NT2RP4001507	C-NT2RP4001507	12278	
NT2RP4001524	C-NT2RP4001524	12279	12280
NT2RP4001547	C-NT2RP4001547	12281	12282
NT2RP4001551	C-NT2RP4001551	12283	12284
NT2RP4001555	C-NT2RP4001555	12285	12286
NT2RP4001567	C-NT2RP4001567	12287	12288
NT2RP4001568	C-NT2RP4001568	12289	12290
NT2RP4001571	C-NT2RP4001571	12291	12292
NT2RP4001574	C-NT2RP4001574	12293	12294
NT2RP4001575	C-NT2RP4001575	12295	12296
NT2RP4001592	C-NT2RP4001592	12297	12298
NT2RP4001610	C-NT2RP4001610	12299	12300
NT2RP4001614	C-NT2RP4001614	12301	12302
NT2RP4001634	C-NT2RP4001634	12303	12304
NT2RP4001638	C-NT2RP4001638	12305	12306
NT2RP4001644	C-NT2RP4001644	12307	12308
NT2RP4001677	C-NT2RP4001677	12309	12310
NT2RP4001679	C-NT2RP4001679	12311	12312
NT2RP4001696	C-NT2RP4001696	12313	12314
NT2RP4001725	C-NT2RP4001725	12315	12316
NT2RP4001730	C-NT2RP4001730	12317	12318
NT2RP4001739	C-NT2RP4001739	12319	12320
NT2RP4001753	C-NT2RP4001753	12321	12322
NT2RP4001760	C-NT2RP4001760	12323	12324
NT2RP4001790	C-NT2RP4001790	12325	12326
NT2RP4001803	C-NT2RP4001803	12327	12328
NT2RP4001822	C-NT2RP4001822	12329	12330
NT2RP4001823	C-NT2RP4001823	12331	12332
NT2RP4001828	C-NT2RP4001828	12333	12334
NT2RP4001838	C-NT2RP4001838	12335	12336
NT2RP4001861	C-NT2RP4001861	12337	12338
NT2RP4001893	C-NT2RP4001893	12339	12340
NT2RP4001896	C-NT2RP4001896	12341	12342
NT2RP4001901	C-NT2RP4001901	12343	12344
NT2RP4001927	C-NT2RP4001927	12345	12346
NT2RP4001938	C-NT2RP4001938	12347	12348
NT2RP4001946	C-NT2RP4001946	12349	12350
NT2RP4001950	C-NT2RP4001950	12351	12352
NT2RP4001953	C-NT2RP4001953	12353	
NT2RP4001966	C-NT2RP4001966	12354	12355
NT2RP4001975	C-NT2RP4001975	12356	12357
NT2RP4002018	C-NT2RP4002018	12358	12359
NT2RP4002052	C-NT2RP4002052	12360	12361
NT2RP4002058	C-NT2RP4002058	12362	12363
NT2RP4002071	C-NT2RP4002071	12364	12365

(940)

特開 2002-191363

1877

1878

NT2RP4002078	C-NT2RP4002078	12366	12367
NT2RP4002081	C-NT2RP4002081	12368	12369
NT2RP4002298	C-NT2RP4002298	12370	12371
NT2RP4002408	C-NT2RP4002408	12372	12373
NT2RP4002791	C-NT2RP4002791	12374	12375
NT2RP4002888	C-NT2RP4002888	12376	12377
NT2RP4002905	C-NT2RP4002905	12378	12379
NT2RP5003461	C-NT2RP5003461	12380	12381
NT2RP5003477	C-NT2RP5003477	12382	12383
NT2RP5003492	C-NT2RP5003492	12384	12385
NT2RP5003500	C-NT2RP5003500	12386	12387
NT2RP5003506	C-NT2RP5003506	12388	12389
NT2RP5003522	C-NT2RP5003522	12390	12391
NT2RP5003524	C-NT2RP5003524	12392	12393
NT2RP5003534	C-NT2RP5003534	12394	12395
OVARC1000006	C-OVARC1000006	12396	12397
OVARC1000013	C-OVARC1000013	12398	12399
OVARC1000014	C-OVARC1000014	12400	12401
OVARC1000035	C-OVARC1000035	12402	
OVARC1000060	C-OVARC1000060	12403	12404
OVARC1000087	C-OVARC1000087	12405	12406
OVARC1000091	C-OVARC1000091	12407	12408
OVARC1000113	C-OVARC1000113	12409	
OVARC1000139	C-OVARC1000139	12410	12411
OVARC1000148	C-OVARC1000148	12412	12413
OVARC1000151	C-OVARC1000151	12414	12415
OVARC1000168	C-OVARC1000168	12416	
OVARC1000209	C-OVARC1000209	12417	12418
OVARC1000212	C-OVARC1000212	12419	12420
OVARC1000241	C-OVARC1000241	12421	12422
OVARC1000288	C-OVARC1000288	12423	12424
OVARC1000304	C-OVARC1000304	12425	12426
OVARC1000309	C-OVARC1000309	12427	12428
OVARC1000321	C-OVARC1000321	12429	12430
OVARC1000326	C-OVARC1000326	12431	12432
OVARC1000335	C-OVARC1000335	12433	12434
OVARC1000347	C-OVARC1000347	12435	
OVARC1000384	C-OVARC1000384	12436	12437
OVARC1000411	C-OVARC1000411	12438	12439
OVARC1000420	C-OVARC1000420	12440	12441
OVARC1000437	C-OVARC1000437	12442	12443
OVARC1000443	C-OVARC1000443	12444	12445
OVARC1000461	C-OVARC1000461	12446	12447
OVARC1000465	C-OVARC1000465	12448	12449
OVARC1000466	C-OVARC1000466	12450	12451
OVARC1000473	C-OVARC1000473	12452	12453
OVARC1000479	C-OVARC1000479	12454	12455
OVARC1000520	C-OVARC1000520	12456	12457
OVARC1000564	C-OVARC1000564	12458	12459
OVARC1000576	C-OVARC1000576	12460	

(941)

特開 2002-191363

1879

1880

OVARC1000588	C-OVARC1000588	12461	12462
OVARC1000605	C-OVARC1000605	12463	12464
OVARC1000640	C-OVARC1000640	12465	12466
OVARC1000649	C-OVARC1000649	12467	12468
OVARC1000661	C-OVARC1000661	12469	12470
OVARC1000771	C-OVARC1000771	12471	12472
OVARC1000959	C-OVARC1000959	12473	12474
OVARC1001034	C-OVARC1001034	12475	12476
OVARC1001038	C-OVARC1001038	12477	12478
OVARC1001065	C-OVARC1001065	12479	12480
OVARC1001162	C-OVARC1001162	12481	
OVARC1001243	C-OVARC1001243	12482	
OVARC1001296	C-OVARC1001296	12483	12484
OVARC1001360	C-OVARC1001360	12485	
OVARC1001381	C-OVARC1001381	12486	12487
OVARC1001425	C-OVARC1001425	12488	12489
PLACE1000005	C-PLACE1000005	12490	
PLACE1000066	C-PLACE1000066	12491	12492
PLACE1000142	C-PLACE1000142	12493	12494
PLACE1000184	C-PLACE1000184	12495	
PLACE1000185	C-PLACE1000185	12496	12497
PLACE1000213	C-PLACE1000213	12498	12499
PLACE1000347	C-PLACE1000347	12500	12501
PLACE1000374	C-PLACE1000374	12502	
PLACE1000380	C-PLACE1000380	12503	12504
PLACE1000383	C-PLACE1000383	12505	12506
PLACE1000401	C-PLACE1000401	12507	12508
PLACE1000406	C-PLACE1000406	12509	12510
PLACE1000420	C-PLACE1000420	12511	12512
PLACE1000435	C-PLACE1000435	12513	
PLACE1000444	C-PLACE1000444	12514	12515
PLACE1000562	C-PLACE1000562	12516	12517
PLACE1000564	C-PLACE1000564	12518	12519
PLACE1000588	C-PLACE1000588	12520	12521
PLACE1000596	C-PLACE1000596	12522	12523
PLACE1000611	C-PLACE1000611	12524	12525
PLACE1000636	C-PLACE1000636	12526	12527
PLACE1000716	C-PLACE1000716	12528	
PLACE1000748	C-PLACE1000748	12529	12530
PLACE1000755	C-PLACE1000755	12531	12532
PLACE1000785	C-PLACE1000785	12533	
PLACE1000798	C-PLACE1000798	12534	
PLACE1000863	C-PLACE1000863	12535	12536
PLACE1000909	C-PLACE1000909	12537	12538
PLACE1000948	C-PLACE1000948	12539	12540
PLACE1000972	C-PLACE1000972	12541	12542
PLACE1000977	C-PLACE1000977	12543	12544
PLACE1001000	C-PLACE1001000	12545	
PLACE1001092	C-PLACE1001092	12546	12547
PLACE1001257	C-PLACE1001257	12548	12549

(942)

特開 2002-191363

1881

1882

PLACE1001383	C-PLACE1001383	12550	12551
PLACE1001387	C-PLACE1001387	12552	12553
PLACE1001399	C-PLACE1001399	12554	
PLACE1001412	C-PLACE1001412	12555	
PLACE1001484	C-PLACE1001484	12556	
PLACE1001503	C-PLACE1001503	12557	12558
PLACE1001570	C-PLACE1001570	12559	12560
PLACE1001610	C-PLACE1001610	12561	12562
PLACE1001692	C-PLACE1001692	12563	12564
PLACE1001729	C-PLACE1001729	12565	12566
PLACE1001739	C-PLACE1001739	12567	12568
PLACE1001781	C-PLACE1001781	12569	12570
PLACE1001810	C-PLACE1001810	12571	
PLACE1001817	C-PLACE1001817	12572	12573
PLACE1001869	C-PLACE1001869	12574	12575
PLACE1001912	C-PLACE1001912	12576	12577
PLACE1001920	C-PLACE1001920	12578	12579
PLACE1001928	C-PLACE1001928	12580	
PLACE1001989	C-PLACE1001989	12581	12582
PLACE1002046	C-PLACE1002046	12583	12584
PLACE1002072	C-PLACE1002072	12585	12586
PLACE1002073	C-PLACE1002073	12587	12588
PLACE1002140	C-PLACE1002140	12589	12590
PLACE1002163	C-PLACE1002163	12591	
PLACE1002170	C-PLACE1002170	12592	
PLACE1002433	C-PLACE1002433	12593	12594
PLACE1002438	C-PLACE1002438	12595	12596
PLACE1002465	C-PLACE1002465	12597	12598
PLACE1002529	C-PLACE1002529	12599	12600
PLACE1002685	C-PLACE1002685	12601	12602
PLACE1002722	C-PLACE1002722	12603	12604
PLACE1002794	C-PLACE1002794	12605	12606
PLACE1002815	C-PLACE1002815	12607	
PLACE1002839	C-PLACE1002839	12608	
PLACE1002851	C-PLACE1002851	12609	12610
PLACE1002941	C-PLACE1002941	12611	12612
PLACE1002996	C-PLACE1002996	12613	12614
PLACE1003045	C-PLACE1003045	12615	12616
PLACE1003092	C-PLACE1003092	12617	12618
PLACE1003100	C-PLACE1003100	12619	12620
PLACE1003108	C-PLACE1003108	12621	
PLACE1003145	C-PLACE1003145	12622	
PLACE1003174	C-PLACE1003174	12623	12624
PLACE1003190	C-PLACE1003190	12625	12626
PLACE1003200	C-PLACE1003200	12627	
PLACE1003296	C-PLACE1003296	12628	12629
PLACE1003302	C-PLACE1003302	12630	12631
PLACE1003334	C-PLACE1003334	12632	12633
PLACE1003342	C-PLACE1003342	12634	12635
PLACE1003353	C-PLACE1003353	12636	12637

(943)

特開 2002-191363

1883

1884

PLACE1003369	C-PLACE1003369	12638	12639
PLACE1003602	C-PLACE1003602	12640	12641
PLACE1003611	C-PLACE1003611	12642	12643
PLACE1003625	C-PLACE1003625	12644	12645
PLACE1003704	C-PLACE1003704	12646	12647
PLACE1003711	C-PLACE1003711	12648	
PLACE1003723	C-PLACE1003723	12649	12650
PLACE1003762	C-PLACE1003762	12651	12652
PLACE1003771	C-PLACE1003771	12653	12654
PLACE1003784	C-PLACE1003784	12655	12656
PLACE1003923	C-PLACE1003923	12657	12658
PLACE1003936	C-PLACE1003936	12659	
PLACE1003968	C-PLACE1003968	12660	12661
PLACE1004104	C-PLACE1004104	12662	12663
PLACE1004114	C-PLACE1004114	12664	
PLACE1004128	C-PLACE1004128	12665	12666
PLACE1004149	C-PLACE1004149	12667	12668
PLACE1004156	C-PLACE1004156	12669	12670
PLACE1004161	C-PLACE1004161	12671	12672
PLACE1004183	C-PLACE1004183	12673	12674
PLACE1004197	C-PLACE1004197	12675	12676
PLACE1004203	C-PLACE1004203	12677	12678
PLACE1004258	C-PLACE1004258	12679	12680
PLACE1004270	C-PLACE1004270	12681	12682
PLACE1004277	C-PLACE1004277	12683	12684
PLACE1004289	C-PLACE1004289	12685	12686
PLACE1004302	C-PLACE1004302	12687	12688
PLACE1004316	C-PLACE1004316	12689	12690
PLACE1004358	C-PLACE1004358	12691	12692
PLACE1004376	C-PLACE1004376	12693	12694
PLACE1004388	C-PLACE1004388	12695	12696
PLACE1004405	C-PLACE1004405	12697	
PLACE1004428	C-PLACE1004428	12698	12699
PLACE1004437	C-PLACE1004437	12700	12701
PLACE1004451	C-PLACE1004451	12702	
PLACE1004460	C-PLACE1004460	12703	12704
PLACE1004473	C-PLACE1004473	12705	12706
PLACE1004510	C-PLACE1004510	12707	12708
PLACE1004516	C-PLACE1004516	12709	
PLACE1004548	C-PLACE1004548	12710	
PLACE1004564	C-PLACE1004564	12711	12712
PLACE1004629	C-PLACE1004629	12713	12714
PLACE1004645	C-PLACE1004645	12715	12716
PLACE1004646	C-PLACE1004646	12717	12718
PLACE1004664	C-PLACE1004664	12719	12720
PLACE1004672	C-PLACE1004672	12721	12722
PLACE1004674	C-PLACE1004674	12723	12724
PLACE1004691	C-PLACE1004691	12725	
PLACE1004722	C-PLACE1004722	12726	12727
PLACE1004736	C-PLACE1004736	12728	12729

1885

1886

PLACE1004740	C-PLACE1004740	12730	
PLACE1004743	C-PLACE1004743	12731	12732
PLACE1004751	C-PLACE1004751	12733	12734
PLACE1004777	C-PLACE1004777	12735	12736
PLACE1004804	C-PLACE1004804	12737	12738
PLACE1004814	C-PLACE1004814	12739	12740
PLACE1004824	C-PLACE1004824	12741	12742
PLACE1004868	C-PLACE1004868	12743	12744
PLACE1004885	C-PLACE1004885	12745	12746
PLACE1004902	C-PLACE1004902	12747	12748
PLACE1004918	C-PLACE1004918	12749	12750
PLACE1004930	C-PLACE1004930	12751	12752
PLACE1004934	C-PLACE1004934	12753	
PLACE1004937	C-PLACE1004937	12754	12755
PLACE1004969	C-PLACE1004969	12756	12757
PLACE1004982	C-PLACE1004982	12758	12759
PLACE1005026	C-PLACE1005026	12760	12761
PLACE1005027	C-PLACE1005027	12762	
PLACE1005046	C-PLACE1005046	12763	
PLACE1005077	C-PLACE1005077	12764	
PLACE1005101	C-PLACE1005101	12765	12766
PLACE1005102	C-PLACE1005102	12767	12768
PLACE1005111	C-PLACE1005111	12769	
PLACE1005181	C-PLACE1005181	12770	
PLACE1005187	C-PLACE1005187	12771	12772
PLACE1005206	C-PLACE1005206	12773	12774
PLACE1005232	C-PLACE1005232	12775	
PLACE1005243	C-PLACE1005243	12776	12777
PLACE1005261	C-PLACE1005261	12778	12779
PLACE1005266	C-PLACE1005266	12780	
PLACE1005277	C-PLACE1005277	12781	12782
PLACE1005287	C-PLACE1005287	12783	12784
PLACE1005305	C-PLACE1005305	12785	12786
PLACE1005308	C-PLACE1005308	12787	12788
PLACE1005313	C-PLACE1005313	12789	12790
PLACE1005327	C-PLACE1005327	12791	12792
PLACE1005335	C-PLACE1005335	12793	12794
PLACE1005373	C-PLACE1005373	12795	12796
PLACE1005374	C-PLACE1005374	12797	
PLACE1005480	C-PLACE1005480	12798	
PLACE1005481	C-PLACE1005481	12799	
PLACE1005494	C-PLACE1005494	12800	
PLACE1005530	C-PLACE1005530	12801	12802
PLACE1005550	C-PLACE1005550	12803	12804
PLACE1005554	C-PLACE1005554	12805	
PLACE1005623	C-PLACE1005623	12806	12807
PLACE1005646	C-PLACE1005646	12808	12809
PLACE1005656	C-PLACE1005656	12810	12811
PLACE1005730	C-PLACE1005730	12812	
PLACE1005755	C-PLACE1005755	12813	12814

1887

1888

PLACE1005763	C-PLACE1005763	12815	12816
PLACE1005803	C-PLACE1005803	12817	12818
PLACE1005804	C-PLACE1005804	12819	12820
PLACE1005851	C-PLACE1005851	12821	12822
PLACE1005921	C-PLACE1005921	12823	12824
PLACE1005923	C-PLACE1005923	12825	12826
PLACE1005925	C-PLACE1005925	12827	12828
PLACE1005934	C-PLACE1005934	12829	12830
PLACE1005936	C-PLACE1005936	12831	12832
PLACE1005951	C-PLACE1005951	12833	12834
PLACE1005953	C-PLACE1005953	12835	12836
PLACE1005955	C-PLACE1005955	12837	12838
PLACE1005966	C-PLACE1005966	12839	12840
PLACE1005990	C-PLACE1005990	12841	12842
PLACE1006011	C-PLACE1006011	12843	12844
PLACE1006040	C-PLACE1006040	12845	12846
PLACE1006119	C-PLACE1006119	12847	
PLACE1006139	C-PLACE1006139	12848	12849
PLACE1006159	C-PLACE1006159	12850	12851
PLACE1006167	C-PLACE1006167	12852	12853
PLACE1006170	C-PLACE1006170	12854	12855
PLACE1006195	C-PLACE1006195	12856	12857
PLACE1006196	C-PLACE1006196	12858	12859
PLACE1006225	C-PLACE1006225	12860	12861
PLACE1006236	C-PLACE1006236	12862	12863
PLACE1006239	C-PLACE1006239	12864	12865
PLACE1006246	C-PLACE1006246	12866	
PLACE1006325	C-PLACE1006325	12867	
PLACE1006335	C-PLACE1006335	12868	12869
PLACE1006357	C-PLACE1006357	12870	
PLACE1006385	C-PLACE1006385	12871	12872
PLACE1006412	C-PLACE1006412	12873	
PLACE1006414	C-PLACE1006414	12874	12875
PLACE1006438	C-PLACE1006438	12876	12877
PLACE1006445	C-PLACE1006445	12878	12879
PLACE1006470	C-PLACE1006470	12880	
PLACE1006482	C-PLACE1006482	12881	12882
PLACE1006488	C-PLACE1006488	12883	12884
PLACE1006492	C-PLACE1006492	12885	12886
PLACE1006531	C-PLACE1006531	12887	12888
PLACE1006552	C-PLACE1006552	12889	12890
PLACE1006598	C-PLACE1006598	12891	
PLACE1006615	C-PLACE1006615	12892	12893
PLACE1006626	C-PLACE1006626	12894	12895
PLACE1006673	C-PLACE1006673	12896	
PLACE1006678	C-PLACE1006678	12897	12898
PLACE1006704	C-PLACE1006704	12899	12900
PLACE1006731	C-PLACE1006731	12901	12902
PLACE1006782	C-PLACE1006782	12903	12904
PLACE1006819	C-PLACE1006819	12905	12906

(946)

特開 2002-191363

1889

1890

PLACE1006829	C-PLACE1006829	12907	12908
PLACE1006883	C-PLACE1006883	12909	12910
PLACE1006901	C-PLACE1006901	12911	12912
PLACE1006917	C-PLACE1006917	12913	12914
PLACE1006932	C-PLACE1006932	12915	
PLACE1006935	C-PLACE1006935	12916	12917
PLACE1006956	C-PLACE1006956	12918	12919
PLACE1006958	C-PLACE1006958	12920	12921
PLACE1006961	C-PLACE1006961	12922	
PLACE1006962	C-PLACE1006962	12923	12924
PLACE1006966	C-PLACE1006966	12925	12926
PLACE1007014	C-PLACE1007014	12927	12928
PLACE1007021	C-PLACE1007021	12929	12930
PLACE1007105	C-PLACE1007105	12931	12932
PLACE1007178	C-PLACE1007178	12933	
PLACE1007226	C-PLACE1007226	12934	12935
PLACE1007238	C-PLACE1007238	12936	12937
PLACE1007239	C-PLACE1007239	12938	12939
PLACE1007242	C-PLACE1007242	12940	
PLACE1007243	C-PLACE1007243	12941	12942
PLACE1007257	C-PLACE1007257	12943	12944
PLACE1007274	C-PLACE1007274	12945	12946
PLACE1007282	C-PLACE1007282	12947	12948
PLACE1007301	C-PLACE1007301	12949	
PLACE1007317	C-PLACE1007317	12950	12951
PLACE1007342	C-PLACE1007342	12952	
PLACE1007346	C-PLACE1007346	12953	12954
PLACE1007367	C-PLACE1007367	12955	
PLACE1007375	C-PLACE1007375	12956	12957
PLACE1007386	C-PLACE1007386	12958	12959
PLACE1007402	C-PLACE1007402	12960	
PLACE1007409	C-PLACE1007409	12961	12962
PLACE1007416	C-PLACE1007416	12963	12964
PLACE1007450	C-PLACE1007450	12965	
PLACE1007452	C-PLACE1007452	12966	
PLACE1007460	C-PLACE1007460	12967	12968
PLACE1007484	C-PLACE1007484	12969	12970
PLACE1007488	C-PLACE1007488	12971	12972
PLACE1007507	C-PLACE1007507	12973	12974
PLACE1007511	C-PLACE1007511	12975	12976
PLACE1007524	C-PLACE1007524	12977	12978
PLACE1007537	C-PLACE1007537	12979	12980
PLACE1007544	C-PLACE1007544	12981	12982
PLACE1007547	C-PLACE1007547	12983	12984
PLACE1007583	C-PLACE1007583	12985	12986
PLACE1007598	C-PLACE1007598	12987	12988
PLACE1007618	C-PLACE1007618	12989	12990
PLACE1007621	C-PLACE1007621	12991	12992
PLACE1007632	C-PLACE1007632	12993	12994
PLACE1007645	C-PLACE1007645	12995	12996

PLACE1007649	C-PLACE1007649	12997	12998
PLACE1007688	C-PLACE1007688	12999	13000
PLACE1007690	C-PLACE1007690	13001	13002
PLACE1007697	C-PLACE1007697	13003	13004
PLACE1007706	C-PLACE1007706	13005	13006
PLACE1007725	C-PLACE1007725	13007	13008
PLACE1007729	C-PLACE1007729	13009	13010
PLACE1007730	C-PLACE1007730	13011	13012
PLACE1007746	C-PLACE1007746	13013	13014
PLACE1007791	C-PLACE1007791	13015	13016
PLACE1007810	C-PLACE1007810	13017	
PLACE1007843	C-PLACE1007843	13018	13019
PLACE1007846	C-PLACE1007846	13020	
PLACE1007858	C-PLACE1007858	13021	13022
PLACE1007897	C-PLACE1007897	13023	13024
PLACE1007946	C-PLACE1007946	13025	13026
PLACE1007954	C-PLACE1007954	13027	13028
PLACE1007955	C-PLACE1007955	13029	13030
PLACE1007958	C-PLACE1007958	13031	13032
PLACE1007969	C-PLACE1007969	13033	13034
PLACE1007990	C-PLACE1007990	13035	13036
PLACE1008000	C-PLACE1008000	13037	13038
PLACE1008002	C-PLACE1008002	13039	
PLACE1008044	C-PLACE1008044	13040	13041
PLACE1008095	C-PLACE1008095	13042	13043
PLACE1008122	C-PLACE1008122	13044	13045
PLACE1008129	C-PLACE1008129	13046	13047
PLACE1008132	C-PLACE1008132	13048	13049
PLACE1008177	C-PLACE1008177	13050	13051
PLACE1008209	C-PLACE1008209	13052	
PLACE1008273	C-PLACE1008273	13053	13054
PLACE1008275	C-PLACE1008275	13055	13056
PLACE1008280	C-PLACE1008280	13057	
PLACE1008309	C-PLACE1008309	13058	13059
PLACE1008329	C-PLACE1008329	13060	
PLACE1008356	C-PLACE1008356	13061	13062
PLACE1008398	C-PLACE1008398	13063	13064
PLACE1008401	C-PLACE1008401	13065	13066
PLACE1008402	C-PLACE1008402	13067	13068
PLACE1008429	C-PLACE1008429	13069	13070
PLACE1008457	C-PLACE1008457	13071	
PLACE1008465	C-PLACE1008465	13072	13073
PLACE1008488	C-PLACE1008488	13074	13075
PLACE1008524	C-PLACE1008524	13076	
PLACE1008531	C-PLACE1008531	13077	13078
PLACE1008532	C-PLACE1008532	13079	13080
PLACE1008533	C-PLACE1008533	13081	13082
PLACE1008568	C-PLACE1008568	13083	13084
PLACE1008603	C-PLACE1008603	13085	13086
PLACE1008621	C-PLACE1008621	13087	

PLACE1008626	C-PLACE1008626	13088	
PLACE1008627	C-PLACE1008627	13089	13090
PLACE1008629	C-PLACE1008629	13091	
PLACE1008650	C-PLACE1008650	13092	13093
PLACE1008693	C-PLACE1008693	13094	13095
PLACE1008696	C-PLACE1008696	13096	
PLACE1008790	C-PLACE1008790	13097	13098
PLACE1008808	C-PLACE1008808	13099	13100
PLACE1008813	C-PLACE1008813	13101	13102
PLACE1008854	C-PLACE1008854	13103	13104
PLACE1008867	C-PLACE1008867	13105	13106
PLACE1008887	C-PLACE1008887	13107	
PLACE1008902	C-PLACE1008902	13108	
PLACE1008925	C-PLACE1008925	13109	
PLACE1009020	C-PLACE1009020	13110	13111
PLACE1009027	C-PLACE1009027	13112	13113
PLACE1009045	C-PLACE1009045	13114	13115
PLACE1009060	C-PLACE1009060	13116	13117
PLACE1009090	C-PLACE1009090	13118	
PLACE1009091	C-PLACE1009091	13119	
PLACE1009094	C-PLACE1009094	13120	13121
PLACE1009099	C-PLACE1009099	13122	13123
PLACE1009110	C-PLACE1009110	13124	
PLACE1009111	C-PLACE1009111	13125	13126
PLACE1009130	C-PLACE1009130	13127	13128
PLACE1009158	C-PLACE1009158	13129	13130
PLACE1009166	C-PLACE1009166	13131	13132
PLACE1009174	C-PLACE1009174	13133	13134
PLACE1009186	C-PLACE1009186	13135	13136
PLACE1009190	C-PLACE1009190	13137	13138
PLACE1009230	C-PLACE1009230	13139	13140
PLACE1009319	C-PLACE1009319	13141	13142
PLACE1009328	C-PLACE1009328	13143	
PLACE1009335	C-PLACE1009335	13144	
PLACE1009338	C-PLACE1009338	13145	13146
PLACE1009368	C-PLACE1009368	13147	13148
PLACE1009375	C-PLACE1009375	13149	13150
PLACE1009388	C-PLACE1009388	13151	
PLACE1009404	C-PLACE1009404	13152	13153
PLACE1009434	C-PLACE1009434	13154	13155
PLACE1009443	C-PLACE1009443	13156	13157
PLACE1009444	C-PLACE1009444	13158	13159
PLACE1009459	C-PLACE1009459	13160	13161
PLACE1009468	C-PLACE1009468	13162	13163
PLACE1009476	C-PLACE1009476	13164	13165
PLACE1009524	C-PLACE1009524	13166	13167
PLACE1009542	C-PLACE1009542	13168	13169
PLACE1009571	C-PLACE1009571	13170	13171
PLACE1009581	C-PLACE1009581	13172	13173
PLACE1009596	C-PLACE1009596	13174	13175

1895

1896

PLACE1009607	C-PLACE1009607	13176	
PLACE1009621	C-PLACE1009621	13177	
PLACE1009622	C-PLACE1009622	13178	13179
PLACE1009659	C-PLACE1009659	13180	
PLACE1009665	C-PLACE1009665	13181	13182
PLACE1009670	C-PLACE1009670	13183	13184
PLACE1009708	C-PLACE1009708	13185	13186
PLACE1009721	C-PLACE1009721	13187	13188
PLACE1009731	C-PLACE1009731	13189	13190
PLACE1009763	C-PLACE1009763	13191	13192
PLACE1009794	C-PLACE1009794	13193	13194
PLACE1009845	C-PLACE1009845	13195	13196
PLACE1009886	C-PLACE1009886	13197	
PLACE1009908	C-PLACE1009908	13198	13199
PLACE1009971	C-PLACE1009971	13200	
PLACE1009992	C-PLACE1009992	13201	13202
PLACE1009995	C-PLACE1009995	13203	13204
PLACE1009997	C-PLACE1009997	13205	13206
PLACE1010023	C-PLACE1010023	13207	13208
PLACE1010031	C-PLACE1010031	13209	13210
PLACE1010053	C-PLACE1010053	13211	13212
PLACE1010074	C-PLACE1010074	13213	13214
PLACE1010076	C-PLACE1010076	13215	
PLACE1010096	C-PLACE1010096	13216	13217
PLACE1010102	C-PLACE1010102	13218	
PLACE1010105	C-PLACE1010105	13219	13220
PLACE1010106	C-PLACE1010106	13221	
PLACE1010134	C-PLACE1010134	13222	13223
PLACE1010148	C-PLACE1010148	13224	13225
PLACE1010194	C-PLACE1010194	13226	13227
PLACE1010202	C-PLACE1010202	13228	13229
PLACE1010261	C-PLACE1010261	13230	13231
PLACE1010274	C-PLACE1010274	13232	13233
PLACE1010293	C-PLACE1010293	13234	
PLACE1010321	C-PLACE1010321	13235	13236
PLACE1010324	C-PLACE1010324	13237	
PLACE1010329	C-PLACE1010329	13238	
PLACE1010362	C-PLACE1010362	13239	13240
PLACE1010364	C-PLACE1010364	13241	
PLACE1010383	C-PLACE1010383	13242	
PLACE1010481	C-PLACE1010481	13243	13244
PLACE1010491	C-PLACE1010491	13245	
PLACE1010492	C-PLACE1010492	13246	13247
PLACE1010522	C-PLACE1010522	13248	13249
PLACE1010529	C-PLACE1010529	13250	13251
PLACE1010547	C-PLACE1010547	13252	13253
PLACE1010599	C-PLACE1010599	13254	13255
PLACE1010616	C-PLACE1010616	13256	
PLACE1010622	C-PLACE1010622	13257	13258
PLACE1010629	C-PLACE1010629	13259	

(950)

特開 2002-191363

1897

1898

PLACE1010630	C-PLACE1010630	13260	
PLACE1010661	C-PLACE1010661	13261	13262
PLACE1010714	C-PLACE1010714	13263	
PLACE1010720	C-PLACE1010720	13264	13265
PLACE1010743	C-PLACE1010743	13266	13267
PLACE1010771	C-PLACE1010771	13268	13269
PLACE1010786	C-PLACE1010786	13270	
PLACE1010800	C-PLACE1010800	13271	13272
PLACE1010811	C-PLACE1010811	13273	13274
PLACE1010870	C-PLACE1010870	13275	
PLACE1010877	C-PLACE1010877	13276	13277
PLACE1010900	C-PLACE1010900	13278	
PLACE2000050	C-PLACE2000050	13279	
PLACE4000522	C-PLACE4000522	13280	13281
PLACE4000590	C-PLACE4000590	13282	13283
PLACE4000638	C-PLACE4000638	13284	13285
PLACE4000650	C-PLACE4000650	13286	13287
Y79AA1001647	C-Y79AA1001647	13288	13289

【0822】

20【表351】

クローン名 配列名	全長塩基配列	全長塩基配列 配列番号	推定アミノ酸配列 配列番号
HEMBA1000012	C-HEMBA1000012	13294	13295
HEMBA1000020	C-HEMBA1000020	13296	13297
HEMBA1000129	C-HEMBA1000129	13298	13299
HEMBA1000201	C-HEMBA1000201	13300	13301
HEMBA1000216	C-HEMBA1000216	13302	13303
HEMBA1000231	C-HEMBA1000231	13304	13305
HEMBA1000264	C-HEMBA1000264	13306	
HEMBA1000280	C-HEMBA1000280	13307	13308
HEMBA1000282	C-HEMBA1000282	13309	
HEMBA1000303	C-HEMBA1000303	13310	13311
HEMBA1000333	C-HEMBA1000333	13312	13313
HEMBA1000351	C-HEMBA1000351	13314	13315
HEMBA1000356	C-HEMBA1000356	13316	13317
HEMBA1000396	C-HEMBA1000396	13318	
HEMBA1000411	C-HEMBA1000411	13319	13320
HEMBA1000442	C-HEMBA1000442	13321	
HEMBA1000456	C-HEMBA1000456	13322	13323
HEMBA1000504	C-HEMBA1000504	13324	
HEMBA1000518	C-HEMBA1000518	13325	13326
HEMBA1000519	C-HEMBA1000519	13327	
HEMBA1000523	C-HEMBA1000523	13328	13329
HEMBA1000542	C-HEMBA1000542	13330	13331
HEMBA1000545	C-HEMBA1000545	13332	
HEMBA1000557	C-HEMBA1000557	13333	
HEMBA1000592	C-HEMBA1000592	13334	13335
HEMBA1000594	C-HEMBA1000594	13336	

(951)

特開 2002-191363

1899

1900

HEMBA1000604	C-HEMBA1000604	13337	
HEMBA1000622	C-HEMBA1000622	13338	
HEMBA1000637	C-HEMBA1000637	13339	13340
HEMBA1000655	C-HEMBA1000655	13341	
HEMBA1000657	C-HEMBA1000657	13342	13343
HEMBA1000749	C-HEMBA1000749	13344	
HEMBA1000769	C-HEMBA1000769	13345	
HEMBA1000773	C-HEMBA1000773	13346	
HEMBA1000774	C-HEMBA1000774	13347	
HEMBA1000822	C-HEMBA1000822	13348	13349
HEMBA1000843	C-HEMBA1000843	13350	
HEMBA1000852	C-HEMBA1000852	13351	
HEMBA1000870	C-HEMBA1000870	13352	13353
HEMBA1000908	C-HEMBA1000908	13354	13355
HEMBA1000934	C-HEMBA1000934	13356	
HEMBA1000972	C-HEMBA1000972	13357	13358
HEMBA1000986	C-HEMBA1000986	13359	
HEMBA1000991	C-HEMBA1000991	13360	13361
HEMBA1001008	C-HEMBA1001008	13362	
HEMBA1001059	C-HEMBA1001059	13363	13364
HEMBA1001094	C-HEMBA1001094	13365	
HEMBA1001302	C-HEMBA1001302	13366	13367
HEMBA1001330	C-HEMBA1001330	13368	
HEMBA1001497	C-HEMBA1001497	13369	
HEMBA1001569	C-HEMBA1001569	13370	
HEMBA1001570	C-HEMBA1001570	13371	
HEMBA1001620	C-HEMBA1001620	13372	13373
HEMBA1001640	C-HEMBA1001640	13374	
HEMBA1001655	C-HEMBA1001655	13375	
HEMBA1001672	C-HEMBA1001672	13376	13377
HEMBA1001711	C-HEMBA1001711	13378	
HEMBA1001723	C-HEMBA1001723	13379	13380
HEMBA1001746	C-HEMBA1001746	13381	13382
HEMBA1001781	C-HEMBA1001781	13383	13384
HEMBA1001804	C-HEMBA1001804	13385	13386
HEMBA1001822	C-HEMBA1001822	13387	13388
HEMBA1001824	C-HEMBA1001824	13389	13390
HEMBA1001866	C-HEMBA1001866	13391	13392
HEMBA1001910	C-HEMBA1001910	13393	
HEMBA1001913	C-HEMBA1001913	13394	13395
HEMBA1001921	C-HEMBA1001921	13396	13397
HEMBA1001939	C-HEMBA1001939	13398	13399
HEMBA1001950	C-HEMBA1001950	13400	13401
HEMBA1001967	C-HEMBA1001967	13402	13403
HEMBA1002035	C-HEMBA1002035	13404	13405
HEMBA1002092	C-HEMBA1002092	13406	13407
HEMBA1002102	C-HEMBA1002102	13408	13409
HEMBA1002150	C-HEMBA1002150	13410	13411
HEMBA1002151	C-HEMBA1002151	13412	13413
HEMBA1002189	C-HEMBA1002189	13414	

(952)

特開 2002-191363

1901

1902

HEMBA1002215	C-HEMBA1002215	13415	13416
HEMBA1002229	C-HEMBA1002229	13417	
HEMBA1002241	C-HEMBA1002241	13418	13419
HEMBA1002341	C-HEMBA1002341	13420	
HEMBA1002417	C-HEMBA1002417	13421	13422
HEMBA1002547	C-HEMBA1002547	13423	13424
HEMBA1002703	C-HEMBA1002703	13425	13426
HEMBA1002779	C-HEMBA1002779	13427	
HEMBA1002816	C-HEMBA1002816	13428	13429
HEMBA1002970	C-HEMBA1002970	13430	
HEMBA1002999	C-HEMBA1002999	13431	13432
HEMBA1003021	C-HEMBA1003021	13433	
HEMBA1003077	C-HEMBA1003077	13434	13435
HEMBA1003079	C-HEMBA1003079	13436	13437
HEMBA1003273	C-HEMBA1003273	13438	
HEMBA1003304	C-HEMBA1003304	13439	
HEMBA1003309	C-HEMBA1003309	13440	
HEMBA1003376	C-HEMBA1003376	13441	
HEMBA1003384	C-HEMBA1003384	13442	
HEMBA1003531	C-HEMBA1003531	13443	
HEMBA1003548	C-HEMBA1003548	13444	
HEMBA1003556	C-HEMBA1003556	13445	13446
HEMBA1003571	C-HEMBA1003571	13447	
HEMBA1003579	C-HEMBA1003579	13448	
HEMBA1003684	C-HEMBA1003684	13449	13450
HEMBA1003692	C-HEMBA1003692	13451	13452
HEMBA1003720	C-HEMBA1003720	13453	
HEMBA1003725	C-HEMBA1003725	13454	
HEMBA1003729	C-HEMBA1003729	13455	13456
HEMBA1003758	C-HEMBA1003758	13457	
HEMBA1003773	C-HEMBA1003773	13458	13459
HEMBA1003783	C-HEMBA1003783	13460	13461
HEMBA1003799	C-HEMBA1003799	13462	
HEMBA1003804	C-HEMBA1003804	13463	13464
HEMBA1003805	C-HEMBA1003805	13465	13466
HEMBA1003836	C-HEMBA1003836	13467	13468
HEMBA1003856	C-HEMBA1003856	13469	
HEMBA1003866	C-HEMBA1003866	13470	13471
HEMBA1003879	C-HEMBA1003879	13472	
HEMBA1003880	C-HEMBA1003880	13473	
HEMBA1003893	C-HEMBA1003893	13474	13475
HEMBA1003908	C-HEMBA1003908	13476	
HEMBA1003937	C-HEMBA1003937	13477	
HEMBA1003942	C-HEMBA1003942	13478	
HEMBA1003958	C-HEMBA1003958	13479	
HEMBA1003976	C-HEMBA1003976	13480	13481
HEMBA1003978	C-HEMBA1003978	13482	
HEMBA1003985	C-HEMBA1003985	13483	
HEMBA1004011	C-HEMBA1004011	13484	
HEMBA1004024	C-HEMBA1004024	13485	

(953)

特開 2002-191363

1903

1904

HEMBA1004038	C-HEMBA1004038	13486	13487
HEMBA1004045	C-HEMBA1004045	13488	
HEMBA1004048	C-HEMBA1004048	13489	13490
HEMBA1004111	C-HEMBA1004111	13491	
HEMBA1004131	C-HEMBA1004131	13492	13493
HEMBA1004138	C-HEMBA1004138	13494	
HEMBA1004143	C-HEMBA1004143	13495	13496
HEMBA1004150	C-HEMBA1004150	13497	
HEMBA1004168	C-HEMBA1004168	13498	13499
HEMBA1004200	C-HEMBA1004200	13500	
HEMBA1004202	C-HEMBA1004202	13501	13502
HEMBA1004203	C-HEMBA1004203	13503	13504
HEMBA1004238	C-HEMBA1004238	13505	
HEMBA1004248	C-HEMBA1004248	13506	
HEMBA1004272	C-HEMBA1004272	13507	
HEMBA1004274	C-HEMBA1004274	13508	13509
HEMBA1004275	C-HEMBA1004275	13510	13511
HEMBA1004286	C-HEMBA1004286	13512	13513
HEMBA1004312	C-HEMBA1004312	13514	
HEMBA1004321	C-HEMBA1004321	13515	13516
HEMBA1004323	C-HEMBA1004323	13517	
HEMBA1004327	C-HEMBA1004327	13518	
HEMBA1004330	C-HEMBA1004330	13519	
HEMBA1004341	C-HEMBA1004341	13520	13521
HEMBA1004366	C-HEMBA1004366	13522	
HEMBA1004372	C-HEMBA1004372	13523	
HEMBA1004389	C-HEMBA1004389	13524	13525
HEMBA1004394	C-HEMBA1004394	13526	
HEMBA1004408	C-HEMBA1004408	13527	13528
HEMBA1004429	C-HEMBA1004429	13529	13530
HEMBA1004460	C-HEMBA1004460	13531	
HEMBA1004461	C-HEMBA1004461	13532	13533
HEMBA1004502	C-HEMBA1004502	13534	
HEMBA1004554	C-HEMBA1004554	13535	
HEMBA1004560	C-HEMBA1004560	13536	13537
HEMBA1004610	C-HEMBA1004610	13538	
HEMBA1004629	C-HEMBA1004629	13539	
HEMBA1004632	C-HEMBA1004632	13540	13541
HEMBA1004637	C-HEMBA1004637	13542	13543
HEMBA1004670	C-HEMBA1004670	13544	13545
HEMBA1004672	C-HEMBA1004672	13546	
HEMBA1004697	C-HEMBA1004697	13547	13548
HEMBA1004711	C-HEMBA1004711	13549	13550
HEMBA1004725	C-HEMBA1004725	13551	13552
HEMBA1004730	C-HEMBA1004730	13553	
HEMBA1004734	C-HEMBA1004734	13554	13555
HEMBA1004751	C-HEMBA1004751	13556	
HEMBA1004752	C-HEMBA1004752	13557	13558
HEMBA1004889	C-HEMBA1004889	13559	13560
HEMBA1004934	C-HEMBA1004934	13561	

(954)

特開 2002-191363

1905

1906

HEMBA1004944	C-HEMBA1004944	13562	
HEMBA1004973	C-HEMBA1004973	13563	13564
HEMBA1004977	C-HEMBA1004977	13565	
HEMBA1005009	C-HEMBA1005009	13566	13567
HEMBA1005083	C-HEMBA1005083	13568	13569
HEMBA1005113	C-HEMBA1005113	13570	
HEMBA1005133	C-HEMBA1005133	13571	
HEMBA1005185	C-HEMBA1005185	13572	13573
HEMBA1005219	C-HEMBA1005219	13574	13575
HEMBA1005252	C-HEMBA1005252	13576	13577
HEMBA1005296	C-HEMBA1005296	13578	13579
HEMBA1005314	C-HEMBA1005314	13580	
HEMBA1005331	C-HEMBA1005331	13581	
HEMBA1005394	C-HEMBA1005394	13582	13583
HEMBA1005403	C-HEMBA1005403	13584	13585
HEMBA1005423	C-HEMBA1005423	13586	13587
HEMBA1005468	C-HEMBA1005468	13588	13589
HEMBA1005469	C-HEMBA1005469	13590	
HEMBA1005474	C-HEMBA1005474	13591	
HEMBA1005517	C-HEMBA1005517	13592	13593
HEMBA1005518	C-HEMBA1005518	13594	
HEMBA1005528	C-HEMBA1005528	13595	13596
HEMBA1005558	C-HEMBA1005558	13597	13598
HEMBA1005576	C-HEMBA1005576	13599	13600
HEMBA1005582	C-HEMBA1005582	13601	13602
HEMBA1005583	C-HEMBA1005583	13603	13604
HEMBA1005595	C-HEMBA1005595	13605	13606
HEMBA1005609	C-HEMBA1005609	13607	
HEMBA1005621	C-HEMBA1005621	13608	13609
HEMBA1005666	C-HEMBA1005666	13610	13611
HEMBA1005680	C-HEMBA1005680	13612	
HEMBA1005685	C-HEMBA1005685	13613	13614
HEMBA1005737	C-HEMBA1005737	13615	13616
HEMBA1005746	C-HEMBA1005746	13617	13618
HEMBA1005755	C-HEMBA1005755	13619	
HEMBA1005813	C-HEMBA1005813	13620	
HEMBA1005822	C-HEMBA1005822	13621	
HEMBA1005834	C-HEMBA1005834	13622	
HEMBA1005884	C-HEMBA1005884	13623	
HEMBA1005891	C-HEMBA1005891	13624	
HEMBA1005909	C-HEMBA1005909	13625	
HEMBA1005911	C-HEMBA1005911	13626	
HEMBA1005931	C-HEMBA1005931	13627	13628
HEMBA1005963	C-HEMBA1005963	13629	13630
HEMBA1005991	C-HEMBA1005991	13631	13632
HEMBA1006005	C-HEMBA1006005	13633	13634
HEMBA1006031	C-HEMBA1006031	13635	13636
HEMBA1006067	C-HEMBA1006067	13637	13638
HEMBA1006081	C-HEMBA1006081	13639	
HEMBA1006091	C-HEMBA1006091	13640	13641

(955)

特開 2002-191363

1907

1908

HEMBA1006100	C-HEMBA1006100	13642	
HEMBA1006108	C-HEMBA1006108	13643	13644
HEMBA1006121	C-HEMBA1006121	13645	13646
HEMBA1006130	C-HEMBA1006130	13647	13648
HEMBA1006155	C-HEMBA1006155	13649	13650
HEMBA1006158	C-HEMBA1006158	13651	
HEMBA1006182	C-HEMBA1006182	13652	
HEMBA1006198	C-HEMBA1006198	13653	13654
HEMBA1006235	C-HEMBA1006235	13655	
HEMBA1006253	C-HEMBA1006253	13656	
HEMBA1006259	C-HEMBA1006259	13657	
HEMBA1006272	C-HEMBA1006272	13658	13659
HEMBA1006278	C-HEMBA1006278	13660	13661
HEMBA1006283	C-HEMBA1006283	13662	13663
HEMBA1006284	C-HEMBA1006284	13664	13665
HEMBA1006291	C-HEMBA1006291	13666	13667
HEMBA1006293	C-HEMBA1006293	13668	13669
HEMBA1006309	C-HEMBA1006309	13670	13671
HEMBA1006349	C-HEMBA1006349	13672	13673
HEMBA1006364	C-HEMBA1006364	13674	
HEMBA1006381	C-HEMBA1006381	13675	
HEMBA1006398	C-HEMBA1006398	13676	13677
HEMBA1006445	C-HEMBA1006445	13678	13679
HEMBA1006483	C-HEMBA1006483	13680	13681
HEMBA1006492	C-HEMBA1006492	13682	13683
HEMBA1006497	C-HEMBA1006497	13684	
HEMBA1006502	C-HEMBA1006502	13685	13686
HEMBA1006507	C-HEMBA1006507	13687	13688
HEMBA1006535	C-HEMBA1006535	13689	
HEMBA1006559	C-HEMBA1006559	13690	13691
HEMBA1006566	C-HEMBA1006566	13692	
HEMBA1006579	C-HEMBA1006579	13693	
HEMBA1006583	C-HEMBA1006583	13694	13695
HEMBA1006612	C-HEMBA1006612	13696	13697
HEMBA1006624	C-HEMBA1006624	13698	13699
HEMBA1006643	C-HEMBA1006643	13700	
HEMBA1006674	C-HEMBA1006674	13701	13702
HEMBA1006682	C-HEMBA1006682	13703	
HEMBA1006708	C-HEMBA1006708	13704	13705
HEMBA1006717	C-HEMBA1006717	13706	13707
HEMBA1006744	C-HEMBA1006744	13708	
HEMBA1006754	C-HEMBA1006754	13709	
HEMBA1006767	C-HEMBA1006767	13710	
HEMBA1006789	C-HEMBA1006789	13711	13712
HEMBA1006832	C-HEMBA1006832	13713	13714
HEMBA1006885	C-HEMBA1006885	13715	13716
HEMBA1006900	C-HEMBA1006900	13717	
HEMBA1006926	C-HEMBA1006926	13718	13719
HEMBA1006941	C-HEMBA1006941	13720	13721
HEMBA1006973	C-HEMBA1006973	13722	13723

(956)

特開 2002-191363

1909

1910

HEMBA1006993	C-HEMBA1006993	13724	
HEMBA1007002	C-HEMBA1007002	13725	13726
HEMBA1007062	C-HEMBA1007062	13727	
HEMBA1007080	C-HEMBA1007080	13728	
HEMBA1007087	C-HEMBA1007087	13729	13730
HEMBA1007112	C-HEMBA1007112	13731	
HEMBA1007194	C-HEMBA1007194	13732	13733
HEMBA1007206	C-HEMBA1007206	13734	
HEMBA1007256	C-HEMBA1007256	13735	
HEMBA1007267	C-HEMBA1007267	13736	13737
HEMBA1007281	C-HEMBA1007281	13738	13739
HEMBA1007300	C-HEMBA1007300	13740	13741
HEMBA1007301	C-HEMBA1007301	13742	13743
HEMBA1007319	C-HEMBA1007319	13744	13745
HEMBA1007320	C-HEMBA1007320	13746	13747
HEMBA1007327	C-HEMBA1007327	13748	
HEMBA1007347	C-HEMBA1007347	13749	
HEMBA1000005	C-HEMBA1000005	13750	
HEMBA1000030	C-HEMBA1000030	13751	13752
HEMBA1000048	C-HEMBA1000048	13753	
HEMBA1000099	C-HEMBA1000099	13754	
HEMBA1000141	C-HEMBA1000141	13755	
HEMBA1000198	C-HEMBA1000198	13756	
HEMBA1000217	C-HEMBA1000217	13757	13758
HEMBA1000218	C-HEMBA1000218	13759	
HEMBA1000274	C-HEMBA1000274	13760	
HEMBA1000312	C-HEMBA1000312	13761	
HEMBA1000402	C-HEMBA1000402	13762	
HEMBA1000420	C-HEMBA1000420	13763	
HEMBA1000480	C-HEMBA1000480	13764	
HEMBA1000530	C-HEMBA1000530	13765	
HEMBA1000550	C-HEMBA1000550	13766	
HEMBA1000556	C-HEMBA1000556	13767	13768
HEMBA1000586	C-HEMBA1000586	13769	
HEMBA1000592	C-HEMBA1000592	13770	13771
HEMBA1000593	C-HEMBA1000593	13772	13773
HEMBA1000649	C-HEMBA1000649	13774	
HEMBA1000693	C-HEMBA1000693	13775	13776
HEMBA1000822	C-HEMBA1000822	13777	
HEMBA1000826	C-HEMBA1000826	13778	
HEMBA1000890	C-HEMBA1000890	13779	
HEMBA1000915	C-HEMBA1000915	13780	13781
HEMBA1001008	C-HEMBA1001008	13782	
HEMBA1001020	C-HEMBA1001020	13783	13784
HEMBA1001051	C-HEMBA1001051	13785	
HEMBA1001112	C-HEMBA1001112	13786	13787
HEMBA1001221	C-HEMBA1001221	13788	
HEMBA1001234	C-HEMBA1001234	13789	13790
HEMBA1001282	C-HEMBA1001282	13791	13792
HEMBA1001302	C-HEMBA1001302	13793	13794

(957)

特開 2002-191363

1911

1912

HEMBB1001335	C-HEMBB1001335	13795	
HEMBB1001337	C-HEMBB1001337	13796	
HEMBB1001356	C-HEMBB1001356	13797	
HEMBB1001364	C-HEMBB1001364	13798	
HEMBB1001366	C-HEMBB1001366	13799	
HEMBB1001367	C-HEMBB1001367	13800	13801
HEMBB1001527	C-HEMBB1001527	13802	
HEMBB1001537	C-HEMBB1001537	13803	
HEMBB1002359	C-HEMBB1002359	13804	
HEMBB1002415	C-HEMBB1002415	13805	
HEMBB1002457	C-HEMBB1002457	13806	13807
HEMBB1002492	C-HEMBB1002492	13808	
HEMBB1002495	C-HEMBB1002495	13809	
HEMBB1002502	C-HEMBB1002502	13810	13811
HEMBB1002550	C-HEMBB1002550	13812	13813
HEMBB1002600	C-HEMBB1002600	13814	13815
HEMBB1002607	C-HEMBB1002607	13816	13817
HEMBB1002684	C-HEMBB1002684	13818	
HEMBB1002692	C-HEMBB1002692	13819	13820
HEMBB1002697	C-HEMBB1002697	13821	13822
HEMBB1002705	C-HEMBB1002705	13823	
MAMMA1000019	C-MAMMA1000019	13824	13825
MAMMA1000020	C-MAMMA1000020	13826	13827
MAMMA1000025	C-MAMMA1000025	13828	
MAMMA1000055	C-MAMMA1000055	13829	13830
MAMMA1000069	C-MAMMA1000069	13831	13832
MAMMA1000084	C-MAMMA1000084	13833	
MAMMA1000139	C-MAMMA1000139	13834	13835
MAMMA1000163	C-MAMMA1000163	13836	13837
MAMMA1000171	C-MAMMA1000171	13838	
MAMMA1000173	C-MAMMA1000173	13839	13840
MAMMA1000277	C-MAMMA1000277	13841	13842
MAMMA1000278	C-MAMMA1000278	13843	13844
MAMMA1000284	C-MAMMA1000284	13845	13846
MAMMA1000309	C-MAMMA1000309	13847	13848
MAMMA1000312	C-MAMMA1000312	13849	
MAMMA1000313	C-MAMMA1000313	13850	
MAMMA1000361	C-MAMMA1000361	13851	
MAMMA1000388	C-MAMMA1000388	13852	13853
MAMMA1000395	C-MAMMA1000395	13854	13855
MAMMA1000410	C-MAMMA1000410	13856	13857
MAMMA1000416	C-MAMMA1000416	13858	13859
MAMMA1000421	C-MAMMA1000421	13860	
MAMMA1000422	C-MAMMA1000422	13861	13862
MAMMA1000468	C-MAMMA1000468	13863	13864
MAMMA1000472	C-MAMMA1000472	13865	13866
MAMMA1000490	C-MAMMA1000490	13867	13868
MAMMA1000524	C-MAMMA1000524	13869	
MAMMA1000567	C-MAMMA1000567	13870	13871
MAMMA1000612	C-MAMMA1000612	13872	13873

(958)

特開 2002-191363

1913

1914

MAMMA1000623	C-MAMMA1000623	13874	13875
MAMMA1000625	C-MAMMA1000625	13876	13877
MAMMA1000664	C-MAMMA1000664	13878	
MAMMA1000670	C-MAMMA1000670	13879	13880
MAMMA1000672	C-MAMMA1000672	13881	13882
MAMMA1000713	C-MAMMA1000713	13883	13884
MAMMA1000731	C-MAMMA1000731	13885	13886
MAMMA1000734	C-MAMMA1000734	13887	13888
MAMMA1000738	C-MAMMA1000738	13889	13890
MAMMA1000746	C-MAMMA1000746	13891	
MAMMA1000775	C-MAMMA1000775	13892	
MAMMA1000824	C-MAMMA1000824	13893	13894
MAMMA1000831	C-MAMMA1000831	13895	
MAMMA1000841	C-MAMMA1000841	13896	13897
MAMMA1000842	C-MAMMA1000842	13898	13899
MAMMA1000843	C-MAMMA1000843	13900	
MAMMA1000856	C-MAMMA1000856	13901	13902
MAMMA1000865	C-MAMMA1000865	13903	
MAMMA1000875	C-MAMMA1000875	13904	
MAMMA1000906	C-MAMMA1000906	13905	
MAMMA1000908	C-MAMMA1000908	13906	13907
MAMMA1000914	C-MAMMA1000914	13908	
MAMMA1000956	C-MAMMA1000956	13909	13910
MAMMA1000968	C-MAMMA1000968	13911	
MAMMA1000979	C-MAMMA1000979	13912	13913
MAMMA1001008	C-MAMMA1001008	13914	13915
MAMMA1001021	C-MAMMA1001021	13916	13917
MAMMA1001041	C-MAMMA1001041	13918	13919
MAMMA1001059	C-MAMMA1001059	13920	13921
MAMMA1001075	C-MAMMA1001075	13922	13923
MAMMA1001078	C-MAMMA1001078	13924	13925
MAMMA1001091	C-MAMMA1001091	13926	13927
MAMMA1001105	C-MAMMA1001105	13928	13929
MAMMA1001110	C-MAMMA1001110	13930	13931
MAMMA1001126	C-MAMMA1001126	13932	
MAMMA1001139	C-MAMMA1001139	13933	13934
MAMMA1001143	C-MAMMA1001143	13935	
MAMMA1001154	C-MAMMA1001154	13936	13937
MAMMA1001181	C-MAMMA1001181	13938	13939
MAMMA1001215	C-MAMMA1001215	13940	
MAMMA1001244	C-MAMMA1001244	13941	
MAMMA1001259	C-MAMMA1001259	13942	13943
MAMMA1001260	C-MAMMA1001260	13944	13945
MAMMA1001343	C-MAMMA1001343	13946	13947
MAMMA1001411	C-MAMMA1001411	13948	13949
MAMMA1001419	C-MAMMA1001419	13950	
MAMMA1001476	C-MAMMA1001476	13951	13952
MAMMA1001510	C-MAMMA1001510	13953	
MAMMA1001522	C-MAMMA1001522	13954	13955
MAMMA1001576	C-MAMMA1001576	13956	13957

(959)

特開 2002-191363

1915

1916

MAMMA1001604	C-MAMMA1001604	13958	
MAMMA1001620	C-MAMMA1001620	13959	13960
MAMMA1001635	C-MAMMA1001635	13961	
MAMMA1001649	C-MAMMA1001649	13962	13963
MAMMA1001686	C-MAMMA1001686	13964	
MAMMA1001692	C-MAMMA1001692	13965	
MAMMA1001743	C-MAMMA1001743	13966	
MAMMA1001754	C-MAMMA1001754	13967	13968
MAMMA1001757	C-MAMMA1001757	13969	
MAMMA1001764	C-MAMMA1001764	13970	
MAMMA1001768	C-MAMMA1001768	13971	13972
MAMMA1001771	C-MAMMA1001771	13973	13974
MAMMA1001790	C-MAMMA1001790	13975	
MAMMA1001837	C-MAMMA1001837	13976	13977
MAMMA1001858	C-MAMMA1001858	13978	
MAMMA1001868	C-MAMMA1001868	13979	13980
MAMMA1001970	C-MAMMA1001970	13981	
MAMMA1002042	C-MAMMA1002042	13982	
MAMMA1002068	C-MAMMA1002068	13983	
MAMMA1002153	C-MAMMA1002153	13984	
MAMMA1002156	C-MAMMA1002156	13985	
MAMMA1002170	C-MAMMA1002170	13986	13987
MAMMA1002174	C-MAMMA1002174	13988	
MAMMA1002209	C-MAMMA1002209	13989	
MAMMA1002219	C-MAMMA1002219	13990	13991
MAMMA1002236	C-MAMMA1002236	13992	13993
MAMMA1002243	C-MAMMA1002243	13994	13995
MAMMA1002268	C-MAMMA1002268	13996	13997
MAMMA1002269	C-MAMMA1002269	13998	
MAMMA1002292	C-MAMMA1002292	13999	14000
MAMMA1002294	C-MAMMA1002294	14001	14002
MAMMA1002297	C-MAMMA1002297	14003	14004
MAMMA1002312	C-MAMMA1002312	14005	
MAMMA1002329	C-MAMMA1002329	14006	14007
MAMMA1002333	C-MAMMA1002333	14008	14009
MAMMA1002351	C-MAMMA1002351	14010	14011
MAMMA1002353	C-MAMMA1002353	14012	
MAMMA1002355	C-MAMMA1002355	14013	
MAMMA1002356	C-MAMMA1002356	14014	
MAMMA1002362	C-MAMMA1002362	14015	
MAMMA1002380	C-MAMMA1002380	14016	14017
MAMMA1002384	C-MAMMA1002384	14018	
MAMMA1002427	C-MAMMA1002427	14019	14020
MAMMA1002470	C-MAMMA1002470	14021	14022
MAMMA1002485	C-MAMMA1002485	14023	14024
MAMMA1002494	C-MAMMA1002494	14025	
MAMMA1002524	C-MAMMA1002524	14026	14027
MAMMA1002530	C-MAMMA1002530	14028	14029
MAMMA1002554	C-MAMMA1002554	14030	
MAMMA1002585	C-MAMMA1002585	14031	14032

(960)

特開 2002-191363

1917

1918

MAMMA1002598	C-MAMMA1002598	14033	14034
MAMMA1002619	C-MAMMA1002619	14035	14036
MAMMA1002655	C-MAMMA1002655	14037	14038
MAMMA1002671	C-MAMMA1002671	14039	14040
MAMMA1002673	C-MAMMA1002673	14041	
MAMMA1002684	C-MAMMA1002684	14042	14043
MAMMA1002711	C-MAMMA1002711	14044	
MAMMA1002769	C-MAMMA1002769	14045	14046
MAMMA1002775	C-MAMMA1002775	14047	14048
MAMMA1002782	C-MAMMA1002782	14049	
MAMMA1002796	C-MAMMA1002796	14050	14051
MAMMA1002807	C-MAMMA1002807	14052	
MAMMA1002838	C-MAMMA1002838	14053	
MAMMA1002842	C-MAMMA1002842	14054	14055
MAMMA1002869	C-MAMMA1002869	14056	14057
MAMMA1002881	C-MAMMA1002881	14058	14059
MAMMA1002886	C-MAMMA1002886	14060	14061
MAMMA1002890	C-MAMMA1002890	14062	14063
MAMMA1002938	C-MAMMA1002938	14064	14065
MAMMA1002964	C-MAMMA1002964	14066	
MAMMA1003011	C-MAMMA1003011	14067	14068
MAMMA1003013	C-MAMMA1003013	14069	14070
MAMMA1003015	C-MAMMA1003015	14071	
MAMMA1003019	C-MAMMA1003019	14072	
MAMMA1003035	C-MAMMA1003035	14073	14074
MAMMA1003039	C-MAMMA1003039	14075	
MAMMA1003044	C-MAMMA1003044	14076	
MAMMA1003049	C-MAMMA1003049	14077	14078
MAMMA1003056	C-MAMMA1003056	14079	14080
MAMMA1003057	C-MAMMA1003057	14081	14082
MAMMA1003066	C-MAMMA1003066	14083	
MAMMA1003099	C-MAMMA1003099	14084	
MAMMA1003104	C-MAMMA1003104	14085	
MAMMA1003113	C-MAMMA1003113	14086	14087
MAMMA1003127	C-MAMMA1003127	14088	14089
MAMMA1003135	C-MAMMA1003135	14090	14091
MAMMA1003146	C-MAMMA1003146	14092	14093
MAMMA1003150	C-MAMMA1003150	14094	14095
MAMMA1003166	C-MAMMA1003166	14096	14097
NT2RM1000032	C-NT2RM1000032	14098	14099
NT2RM1000035	C-NT2RM1000035	14100	14101
NT2RM1000039	C-NT2RM1000039	14102	14103
NT2RM1000055	C-NT2RM1000055	14104	14105
NT2RM1000059	C-NT2RM1000059	14106	14107
NT2RM1000062	C-NT2RM1000062	14108	14109
NT2RM1000118	C-NT2RM1000118	14110	14111
NT2RM1000119	C-NT2RM1000119	14112	14113
NT2RM1000127	C-NT2RM1000127	14114	14115
NT2RM1000131	C-NT2RM1000131	14116	14117
NT2RM1000132	C-NT2RM1000132	14118	14119

1919

1920

NT2RM1000153	C-NT2RM1000153	14120	14121
NT2RM1000186	C-NT2RM1000186	14122	14123
NT2RM1000187	C-NT2RM1000187	14124	14125
NT2RM1000199	C-NT2RM1000199	14126	14127
NT2RM1000244	C-NT2RM1000244	14128	
NT2RM1000252	C-NT2RM1000252	14129	14130
NT2RM1000256	C-NT2RM1000256	14131	14132
NT2RM1000260	C-NT2RM1000260	14133	14134
NT2RM1000271	C-NT2RM1000271	14135	14136
NT2RM1000300	C-NT2RM1000300	14137	14138
NT2RM1000314	C-NT2RM1000314	14139	14140
NT2RM1000354	C-NT2RM1000354	14141	14142
NT2RM1000355	C-NT2RM1000355	14143	14144
NT2RM1000365	C-NT2RM1000365	14145	
NT2RM1000377	C-NT2RM1000377	14146	14147
NT2RM1000388	C-NT2RM1000388	14148	14149
NT2RM1000399	C-NT2RM1000399	14150	14151
NT2RM1000430	C-NT2RM1000430	14152	14153
NT2RM1000555	C-NT2RM1000555	14154	14155
NT2RM1000563	C-NT2RM1000563	14156	14157
NT2RM1000648	C-NT2RM1000648	14158	14159
NT2RM1000661	C-NT2RM1000661	14160	14161
NT2RM1000666	C-NT2RM1000666	14162	14163
NT2RM1000672	C-NT2RM1000672	14164	
NT2RM1000691	C-NT2RM1000691	14165	14166
NT2RM1000699	C-NT2RM1000699	14167	
NT2RM1000741	C-NT2RM1000741	14168	
NT2RM1000742	C-NT2RM1000742	14169	14170
NT2RM1000746	C-NT2RM1000746	14171	14172
NT2RM1000770	C-NT2RM1000770	14173	14174
NT2RM1000772	C-NT2RM1000772	14175	14176
NT2RM1000780	C-NT2RM1000780	14177	14178
NT2RM1000800	C-NT2RM1000800	14179	14180
NT2RM1000802	C-NT2RM1000802	14181	14182
NT2RM1000811	C-NT2RM1000811	14183	14184
NT2RM1000826	C-NT2RM1000826	14185	14186
NT2RM1000829	C-NT2RM1000829	14187	
NT2RM1000850	C-NT2RM1000850	14188	14189
NT2RM1000852	C-NT2RM1000852	14190	14191
NT2RM1000857	C-NT2RM1000857	14192	14193
NT2RM1000874	C-NT2RM1000874	14194	14195
NT2RM1000882	C-NT2RM1000882	14196	14197
NT2RM1000885	C-NT2RM1000885	14198	14199
NT2RM1000894	C-NT2RM1000894	14200	14201
NT2RM1000898	C-NT2RM1000898	14202	14203
NT2RM1000905	C-NT2RM1000905	14204	14205
NT2RM1000924	C-NT2RM1000924	14206	14207
NT2RM1000927	C-NT2RM1000927	14208	14209
NT2RM1000962	C-NT2RM1000962	14210	14211
NT2RM1000978	C-NT2RM1000978	14212	14213

1921

1922

NT2RM1001003	C-NT2RM1001003	14214	14215
NT2RM1001043	C-NT2RM1001043	14216	
NT2RM1001066	C-NT2RM1001066	14217	14218
NT2RM1001072	C-NT2RM1001072	14219	14220
NT2RM1001085	C-NT2RM1001085	14221	14222
NT2RM1001102	C-NT2RM1001102	14223	14224
NT2RM1001105	C-NT2RM1001105	14225	14226
NT2RM1001139	C-NT2RM1001139	14227	14228
NT2RM2000420	C-NT2RM2000420	14229	14230
NT2RM2000566	C-NT2RM2000566	14231	14232
NT2RM2000609	C-NT2RM2000609	14233	14234
NT2RM2000612	C-NT2RM2000612	14235	14236
NT2RM2000735	C-NT2RM2000735	14237	14238
NT2RM2001588	C-NT2RM2001588	14239	14240
NT2RM2001605	C-NT2RM2001605	14241	14242
NT2RM2001613	C-NT2RM2001613	14243	14244
NT2RM2001632	C-NT2RM2001632	14245	14246
NT2RM2001648	C-NT2RM2001648	14247	14248
NT2RM2001652	C-NT2RM2001652	14249	14250
NT2RM2001659	C-NT2RM2001659	14251	14252
NT2RM2001664	C-NT2RM2001664	14253	14254
NT2RM2001668	C-NT2RM2001668	14255	14256
NT2RM2001671	C-NT2RM2001671	14257	14258
NT2RM2001675	C-NT2RM2001675	14259	14260
NT2RM2001681	C-NT2RM2001681	14261	14262
NT2RM2001688	C-NT2RM2001688	14263	14264
NT2RM2001695	C-NT2RM2001695	14265	
NT2RM2001696	C-NT2RM2001696	14266	14267
NT2RM2001698	C-NT2RM2001698	14268	14269
NT2RM2001700	C-NT2RM2001700	14270	14271
NT2RM2001716	C-NT2RM2001716	14272	14273
NT2RM2001723	C-NT2RM2001723	14274	14275
NT2RM2001730	C-NT2RM2001730	14276	14277
NT2RM2001743	C-NT2RM2001743	14278	14279
NT2RM2001753	C-NT2RM2001753	14280	14281
NT2RM2001760	C-NT2RM2001760	14282	14283
NT2RM2001768	C-NT2RM2001768	14284	14285
NT2RM2001771	C-NT2RM2001771	14286	
NT2RM2001782	C-NT2RM2001782	14287	14288
NT2RM2001784	C-NT2RM2001784	14289	14290
NT2RM2001785	C-NT2RM2001785	14291	14292
NT2RM2001813	C-NT2RM2001813	14293	14294
NT2RM2001823	C-NT2RM2001823	14295	14296
NT2RM2001839	C-NT2RM2001839	14297	14298
NT2RM2001840	C-NT2RM2001840	14299	
NT2RM2001855	C-NT2RM2001855	14300	14301
NT2RM2001867	C-NT2RM2001867	14302	14303
NT2RM2001879	C-NT2RM2001879	14304	
NT2RM2001983	C-NT2RM2001983	14305	14306
NT2RM2002145	C-NT2RM2002145	14307	14308

(963)

特開 2002-191363

1923

1924

NT2RM4000027	C-NT2RM4000027	14309	14310
NT2RM4000030	C-NT2RM4000030	14311	14312
NT2RM4000046	C-NT2RM4000046	14313	14314
NT2RM4000155	C-NT2RM4000155	14315	14316
NT2RM4000156	C-NT2RM4000156	14317	14318
NT2RM4000167	C-NT2RM4000167	14319	14320
NT2RM4000199	C-NT2RM4000199	14321	14322
NT2RM4000200	C-NT2RM4000200	14323	14324
NT2RM4000202	C-NT2RM4000202	14325	14326
NT2RM4000233	C-NT2RM4000233	14327	14328
NT2RM4000244	C-NT2RM4000244	14329	
NT2RM4000251	C-NT2RM4000251	14330	14331
NT2RM4000265	C-NT2RM4000265	14332	
NT2RM4000324	C-NT2RM4000324	14333	14334
NT2RM4000327	C-NT2RM4000327	14335	
NT2RM4000356	C-NT2RM4000356	14336	14337
NT2RM4000425	C-NT2RM4000425	14338	
NT2RM4000433	C-NT2RM4000433	14339	14340
NT2RM4000514	C-NT2RM4000514	14341	14342
NT2RM4000531	C-NT2RM4000531	14343	
NT2RM4000532	C-NT2RM4000532	14344	
NT2RM4000534	C-NT2RM4000534	14345	14346
NT2RM4000603	C-NT2RM4000603	14347	14348
NT2RM4000611	C-NT2RM4000611	14349	14350
NT2RM4000616	C-NT2RM4000616	14351	14352
NT2RM4000674	C-NT2RM4000674	14353	14354
NT2RM4000689	C-NT2RM4000689	14355	14356
NT2RM4000698	C-NT2RM4000698	14357	14358
NT2RM4000700	C-NT2RM4000700	14359	14360
NT2RM4000712	C-NT2RM4000712	14361	14362
NT2RM4000717	C-NT2RM4000717	14363	14364
NT2RM4000733	C-NT2RM4000733	14365	14366
NT2RM4000734	C-NT2RM4000734	14367	14368
NT2RM4000741	C-NT2RM4000741	14369	14370
NT2RM4000751	C-NT2RM4000751	14371	14372
NT2RM4000764	C-NT2RM4000764	14373	14374
NT2RM4000778	C-NT2RM4000778	14375	14376
NT2RM4000787	C-NT2RM4000787	14377	
NT2RM4000790	C-NT2RM4000790	14378	14379
NT2RM4000795	C-NT2RM4000795	14380	14381
NT2RM4000796	C-NT2RM4000796	14382	14383
NT2RM4000798	C-NT2RM4000798	14384	14385
NT2RM4000813	C-NT2RM4000813	14386	14387
NT2RM4000820	C-NT2RM4000820	14388	14389
NT2RM4000833	C-NT2RM4000833	14390	14391
NT2RM4000848	C-NT2RM4000848	14392	14393
NT2RM4000852	C-NT2RM4000852	14394	
NT2RM4000855	C-NT2RM4000855	14395	14396
NT2RM4000887	C-NT2RM4000887	14397	14398
NT2RM4000895	C-NT2RM4000895	14399	

1925

1926

NT2RM4000950	C-NT2RM4000950	14400	14401
NT2RM4000979	C-NT2RM4000979	14402	
NT2RM4001002	C-NT2RM4001002	14403	14404
NT2RM4001032	C-NT2RM4001032	14405	14406
NT2RM4001047	C-NT2RM4001047	14407	14408
NT2RM4001054	C-NT2RM4001054	14409	14410
NT2RM4001084	C-NT2RM4001084	14411	14412
NT2RM4001116	C-NT2RM4001116	14413	14414
NT2RM4001140	C-NT2RM4001140	14415	14416
NT2RM4001151	C-NT2RM4001151	14417	14418
NT2RM4001155	C-NT2RM4001155	14419	14420
NT2RM4001160	C-NT2RM4001160	14421	
NT2RM4001187	C-NT2RM4001187	14422	14423
NT2RM4001191	C-NT2RM4001191	14424	14425
NT2RM4001200	C-NT2RM4001200	14426	14427
NT2RM4001203	C-NT2RM4001203	14428	14429
NT2RM4001204	C-NT2RM4001204	14430	14431
NT2RM4001217	C-NT2RM4001217	14432	14433
NT2RM4001256	C-NT2RM4001256	14434	14435
NT2RM4001258	C-NT2RM4001258	14436	14437
NT2RM4001309	C-NT2RM4001309	14438	
NT2RM4001313	C-NT2RM4001313	14439	14440
NT2RM4001316	C-NT2RM4001316	14441	14442
NT2RM4001320	C-NT2RM4001320	14443	14444
NT2RM4001340	C-NT2RM4001340	14445	14446
NT2RM4001344	C-NT2RM4001344	14447	14448
NT2RM4001347	C-NT2RM4001347	14449	14450
NT2RM4001371	C-NT2RM4001371	14451	14452
NT2RM4001382	C-NT2RM4001382	14453	14454
NT2RM4001384	C-NT2RM4001384	14455	14456
NT2RM4001410	C-NT2RM4001410	14457	14458
NT2RM4001411	C-NT2RM4001411	14459	14460
NT2RM4001412	C-NT2RM4001412	14461	14462
NT2RM4001414	C-NT2RM4001414	14463	
NT2RM4001437	C-NT2RM4001437	14464	
NT2RM4001444	C-NT2RM4001444	14465	14466
NT2RM4001454	C-NT2RM4001454	14467	14468
NT2RM4001455	C-NT2RM4001455	14469	14470
NT2RM4001483	C-NT2RM4001483	14471	14472
NT2RM4001489	C-NT2RM4001489	14473	14474
NT2RM4001522	C-NT2RM4001522	14475	14476
NT2RM4001557	C-NT2RM4001557	14477	14478
NT2RM4001565	C-NT2RM4001565	14479	14480
NT2RM4001566	C-NT2RM4001566	14481	14482
NT2RM4001582	C-NT2RM4001582	14483	14484
NT2RM4001592	C-NT2RM4001592	14485	14486
NT2RM4001594	C-NT2RM4001594	14487	14488
NT2RM4001597	C-NT2RM4001597	14489	14490
NT2RM4001611	C-NT2RM4001611	14491	14492
NT2RM4001629	C-NT2RM4001629	14493	14494

(965)

特開 2002-191363

1927

1928

NT2RM4001650	C-NT2RM4001650	14495	14496
NT2RM4001662	C-NT2RM4001662	14497	14498
NT2RM4001666	C-NT2RM4001666	14499	14500
NT2RM4001682	C-NT2RM4001682	14501	14502
NT2RM4001710	C-NT2RM4001710	14503	
NT2RM4001714	C-NT2RM4001714	14504	14505
NT2RM4001715	C-NT2RM4001715	14506	14507
NT2RM4001731	C-NT2RM4001731	14508	14509
NT2RM4001746	C-NT2RM4001746	14510	14511
NT2RM4001754	C-NT2RM4001754	14512	
NT2RM4001758	C-NT2RM4001758	14513	14514
NT2RM4001783	C-NT2RM4001783	14515	14516
NT2RM4001810	C-NT2RM4001810	14517	14518
NT2RM4001813	C-NT2RM4001813	14519	14520
NT2RM4001823	C-NT2RM4001823	14521	14522
NT2RM4001828	C-NT2RM4001828	14523	14524
NT2RM4001836	C-NT2RM4001836	14525	14526
NT2RM4001841	C-NT2RM4001841	14527	
NT2RM4001842	C-NT2RM4001842	14528	
NT2RM4001856	C-NT2RM4001856	14529	14530
NT2RM4001858	C-NT2RM4001858	14531	14532
NT2RM4001865	C-NT2RM4001865	14533	14534
NT2RM4001876	C-NT2RM4001876	14535	14536
NT2RM4001880	C-NT2RM4001880	14537	14538
NT2RM4001922	C-NT2RM4001922	14539	
NT2RM4001930	C-NT2RM4001930	14540	14541
NT2RM4001940	C-NT2RM4001940	14542	14543
NT2RM4001953	C-NT2RM4001953	14544	
NT2RM4001965	C-NT2RM4001965	14545	14546
NT2RM4001969	C-NT2RM4001969	14547	14548
NT2RM4001979	C-NT2RM4001979	14549	14550
NT2RM4001984	C-NT2RM4001984	14551	
NT2RM4001987	C-NT2RM4001987	14552	14553
NT2RM4002013	C-NT2RM4002013	14554	14555
NT2RM4002018	C-NT2RM4002018	14556	14557
NT2RM4002034	C-NT2RM4002034	14558	14559
NT2RM4002044	C-NT2RM4002044	14560	14561
NT2RM4002054	C-NT2RM4002054	14562	14563
NT2RM4002063	C-NT2RM4002063	14564	14565
NT2RM4002066	C-NT2RM4002066	14566	14567
NT2RM4002075	C-NT2RM4002075	14568	14569
NT2RM4002128	C-NT2RM4002128	14570	14571
NT2RM4002140	C-NT2RM4002140	14572	14573
NT2RM4002145	C-NT2RM4002145	14574	14575
NT2RM4002161	C-NT2RM4002161	14576	14577
NT2RM4002174	C-NT2RM4002174	14578	14579
NT2RM4002189	C-NT2RM4002189	14580	14581
NT2RM4002205	C-NT2RM4002205	14582	14583
NT2RM4002213	C-NT2RM4002213	14584	14585
NT2RM4002226	C-NT2RM4002226	14586	14587

(966)

特開 2002-191363

1929

1930

NT2RM4002251	C-NT2RM4002251	14588	14589
NT2RM4002256	C-NT2RM4002256	14590	14591
NT2RM4002266	C-NT2RM4002266	14592	14593
NT2RM4002281	C-NT2RM4002281	14594	
NT2RM4002287	C-NT2RM4002287	14595	14596
NT2RM4002294	C-NT2RM4002294	14597	14598
NT2RM4002301	C-NT2RM4002301	14599	14600
NT2RM4002323	C-NT2RM4002323	14601	14602
NT2RM4002339	C-NT2RM4002339	14603	14604
NT2RM4002344	C-NT2RM4002344	14605	14606
NT2RM4002373	C-NT2RM4002373	14607	14608
NT2RM4002374	C-NT2RM4002374	14609	
NT2RM4002383	C-NT2RM4002383	14610	
NT2RM4002409	C-NT2RM4002409	14611	14612
NT2RM4002438	C-NT2RM4002438	14613	14614
NT2RM4002446	C-NT2RM4002446	14615	14616
NT2RM4002452	C-NT2RM4002452	14617	14618
NT2RM4002457	C-NT2RM4002457	14619	
NT2RM4002460	C-NT2RM4002460	14620	14621
NT2RM4002493	C-NT2RM4002493	14622	14623
NT2RM4002527	C-NT2RM4002527	14624	14625
NT2RM4002532	C-NT2RM4002532	14626	14627
NT2RM4002558	C-NT2RM4002558	14628	14629
NT2RM4002567	C-NT2RM4002567	14630	14631
NT2RM4002593	C-NT2RM4002593	14632	
NT2RM4002594	C-NT2RM4002594	14633	14634
NT2RM4002623	C-NT2RM4002623	14635	14636
NT2RP1000324	C-NT2RP1000324	14637	14638
NT2RP1000363	C-NT2RP1000363	14639	14640
NT2RP1000418	C-NT2RP1000418	14641	14642
NT2RP1000513	C-NT2RP1000513	14643	
NT2RP1000721	C-NT2RP1000721	14644	14645
NT2RP1000730	C-NT2RP1000730	14646	14647
NT2RP1000767	C-NT2RP1000767	14648	14649
NT2RP1000836	C-NT2RP1000836	14650	14651
NT2RP1000902	C-NT2RP1000902	14652	14653
NT2RP1000943	C-NT2RP1000943	14654	14655
NT2RP1001033	C-NT2RP1001033	14656	14657
NT2RP1001073	C-NT2RP1001073	14658	14659
NT2RP1001199	C-NT2RP1001199	14660	14661
NT2RP1001248	C-NT2RP1001248	14662	14663
NT2RP1001253	C-NT2RP1001253	14664	14665
NT2RP1001286	C-NT2RP1001286	14666	14667
NT2RP1001294	C-NT2RP1001294	14668	14669
NT2RP1001302	C-NT2RP1001302	14670	14671
NT2RP1001310	C-NT2RP1001310	14672	14673
NT2RP1001361	C-NT2RP1001361	14674	14675
NT2RP1001385	C-NT2RP1001385	14676	14677
NT2RP1001432	C-NT2RP1001432	14678	
NT2RP2000040	C-NT2RP2000040	14679	14680

(967)

特開 2002-191363

1931

1932

NT2RP2000076	C-NT2RP2000076	14681	14682
NT2RP2000098	C-NT2RP2000098	14683	14684
NT2RP2000108	C-NT2RP2000108	14685	
NT2RP2000257	C-NT2RP2000257	14686	14687
NT2RP2000258	C-NT2RP2000258	14688	14689
NT2RP2000289	C-NT2RP2000289	14690	14691
NT2RP2000327	C-NT2RP2000327	14692	
NT2RP2000337	C-NT2RP2000337	14693	14694
NT2RP2000420	C-NT2RP2000420	14695	14696
NT2RP2000459	C-NT2RP2000459	14697	14698
NT2RP2000498	C-NT2RP2000498	14699	
NT2RP2000758	C-NT2RP2000758	14700	14701
NT2RP2001137	C-NT2RP2001137	14702	14703
NT2RP2001149	C-NT2RP2001149	14704	14705
NT2RP2001168	C-NT2RP2001168	14706	14707
NT2RP2001173	C-NT2RP2001173	14708	14709
NT2RP2001174	C-NT2RP2001174	14710	14711
NT2RP2001196	C-NT2RP2001196	14712	
NT2RP2001226	C-NT2RP2001226	14713	14714
NT2RP2001268	C-NT2RP2001268	14715	14716
NT2RP2001290	C-NT2RP2001290	14717	14718
NT2RP2001295	C-NT2RP2001295	14719	14720
NT2RP2001312	C-NT2RP2001312	14721	
NT2RP2001327	C-NT2RP2001327	14722	14723
NT2RP2001328	C-NT2RP2001328	14724	
NT2RP2001366	C-NT2RP2001366	14725	14726
NT2RP2001378	C-NT2RP2001378	14727	14728
NT2RP2001392	C-NT2RP2001392	14729	14730
NT2RP2001394	C-NT2RP2001394	14731	
NT2RP2001420	C-NT2RP2001420	14732	14733
NT2RP2001450	C-NT2RP2001450	14734	
NT2RP2001467	C-NT2RP2001467	14735	14736
NT2RP2001506	C-NT2RP2001506	14737	
NT2RP2001511	C-NT2RP2001511	14738	14739
NT2RP2001520	C-NT2RP2001520	14740	14741
NT2RP2001536	C-NT2RP2001536	14742	14743
NT2RP2001560	C-NT2RP2001560	14744	14745
NT2RP2001576	C-NT2RP2001576	14746	14747
NT2RP2001581	C-NT2RP2001581	14748	14749
NT2RP2001597	C-NT2RP2001597	14750	14751
NT2RP2001628	C-NT2RP2001628	14752	14753
NT2RP2001663	C-NT2RP2001663	14754	
NT2RP2001748	C-NT2RP2001748	14755	14756
NT2RP2001813	C-NT2RP2001813	14757	
NT2RP2001883	C-NT2RP2001883	14758	14759
NT2RP2001900	C-NT2RP2001900	14760	14761
NT2RP2001947	C-NT2RP2001947	14762	14763
NT2RP2001985	C-NT2RP2001985	14764	14765
NT2RP2001991	C-NT2RP2001991	14766	14767
NT2RP2002025	C-NT2RP2002025	14768	14769

(968)

特開 2002-191363

1933

1934

NT2RP2002058	C-NT2RP2002058	14770	14771
NT2RP2002076	C-NT2RP2002076	14772	14773
NT2RP2002078	C-NT2RP2002078	14774	14775
NT2RP2002079	C-NT2RP2002079	14776	14777
NT2RP2002099	C-NT2RP2002099	14778	14779
NT2RP2002185	C-NT2RP2002185	14780	14781
NT2RP2002193	C-NT2RP2002193	14782	14783
NT2RP2002231	C-NT2RP2002231	14784	14785
NT2RP2002235	C-NT2RP2002235	14786	14787
NT2RP2002252	C-NT2RP2002252	14788	14789
NT2RP2002292	C-NT2RP2002292	14790	14791
NT2RP2002408	C-NT2RP2002408	14792	14793
NT2RP2002442	C-NT2RP2002442	14794	14795
NT2RP2002464	C-NT2RP2002464	14796	14797
NT2RP2002498	C-NT2RP2002498	14798	14799
NT2RP2002503	C-NT2RP2002503	14800	14801
NT2RP2002520	C-NT2RP2002520	14802	14803
NT2RP2002549	C-NT2RP2002549	14804	14805
NT2RP2002609	C-NT2RP2002609	14806	14807
NT2RP2002706	C-NT2RP2002706	14808	
NT2RP2002710	C-NT2RP2002710	14809	14810
NT2RP2002800	C-NT2RP2002800	14811	
NT2RP2002880	C-NT2RP2002880	14812	14813
NT2RP2002891	C-NT2RP2002891	14814	14815
NT2RP2002929	C-NT2RP2002929	14816	14817
NT2RP2002939	C-NT2RP2002939	14818	14819
NT2RP2002993	C-NT2RP2002993	14820	14821
NT2RP2003034	C-NT2RP2003034	14822	
NT2RP2003099	C-NT2RP2003099	14823	
NT2RP2003137	C-NT2RP2003137	14824	14825
NT2RP2003157	C-NT2RP2003157	14826	14827
NT2RP2003158	C-NT2RP2003158	14828	14829
NT2RP2003165	C-NT2RP2003165	14830	
NT2RP2003243	C-NT2RP2003243	14831	14832
NT2RP2003277	C-NT2RP2003277	14833	14834
NT2RP2003286	C-NT2RP2003286	14835	14836
NT2RP2003297	C-NT2RP2003297	14837	14838
NT2RP2003307	C-NT2RP2003307	14839	14840
NT2RP2003308	C-NT2RP2003308	14841	14842
NT2RP2003347	C-NT2RP2003347	14843	14844
NT2RP2003391	C-NT2RP2003391	14845	14846
NT2RP2003393	C-NT2RP2003393	14847	14848
NT2RP2003445	C-NT2RP2003445	14849	
NT2RP2003466	C-NT2RP2003466	14850	14851
NT2RP2003480	C-NT2RP2003480	14852	14853
NT2RP2003506	C-NT2RP2003506	14854	14855
NT2RP2003511	C-NT2RP2003511	14856	14857
NT2RP2003513	C-NT2RP2003513	14858	14859
NT2RP2003567	C-NT2RP2003567	14860	14861
NT2RP2003604	C-NT2RP2003604	14862	14863

(969)

特開 2002-191363

1935

1936

NT2RP2003691	C-NT2RP2003691	14864	
NT2RP2003713	C-NT2RP2003713	14865	14866
NT2RP2003760	C-NT2RP2003760	14867	14868
NT2RP2003764	C-NT2RP2003764	14869	14870
NT2RP2003769	C-NT2RP2003769	14871	
NT2RP2003777	C-NT2RP2003777	14872	14873
NT2RP2003840	C-NT2RP2003840	14874	14875
NT2RP2003857	C-NT2RP2003857	14876	14877
NT2RP2003981	C-NT2RP2003981	14878	14879
NT2RP2003984	C-NT2RP2003984	14880	14881
NT2RP2004041	C-NT2RP2004041	14882	14883
NT2RP2004066	C-NT2RP2004066	14884	14885
NT2RP2004081	C-NT2RP2004081	14886	14887
NT2RP2004124	C-NT2RP2004124	14888	14889
NT2RP2004152	C-NT2RP2004152	14890	14891
NT2RP2004165	C-NT2RP2004165	14892	14893
NT2RP2004187	C-NT2RP2004187	14894	14895
NT2RP2004239	C-NT2RP2004239	14896	14897
NT2RP2004245	C-NT2RP2004245	14898	14899
NT2RP2004364	C-NT2RP2004364	14900	14901
NT2RP2004365	C-NT2RP2004365	14902	14903
NT2RP2004366	C-NT2RP2004366	14904	14905
NT2RP2004373	C-NT2RP2004373	14906	14907
NT2RP2004476	C-NT2RP2004476	14908	14909
NT2RP2004551	C-NT2RP2004551	14910	14911
NT2RP2004568	C-NT2RP2004568	14912	14913
NT2RP2004600	C-NT2RP2004600	14914	14915
NT2RP2004664	C-NT2RP2004664	14916	14917
NT2RP2004743	C-NT2RP2004743	14918	
NT2RP2004768	C-NT2RP2004768	14919	14920
NT2RP2004816	C-NT2RP2004816	14921	14922
NT2RP2004861	C-NT2RP2004861	14923	
NT2RP2004897	C-NT2RP2004897	14924	
NT2RP2004933	C-NT2RP2004933	14925	14926
NT2RP2004978	C-NT2RP2004978	14927	14928
NT2RP2005038	C-NT2RP2005038	14929	14930
NT2RP2005162	C-NT2RP2005162	14931	14932
NT2RP2005204	C-NT2RP2005204	14933	14934
NT2RP2005227	C-NT2RP2005227	14935	14936
NT2RP2005287	C-NT2RP2005287	14937	
NT2RP2005288	C-NT2RP2005288	14938	14939
NT2RP2005490	C-NT2RP2005490	14940	14941
NT2RP2005539	C-NT2RP2005539	14942	14943
NT2RP2005605	C-NT2RP2005605	14944	14945
NT2RP2005722	C-NT2RP2005722	14946	14947
NT2RP2005732	C-NT2RP2005732	14948	14949
NT2RP2005784	C-NT2RP2005784	14950	14951
NT2RP2005812	C-NT2RP2005812	14952	14953
NT2RP2005859	C-NT2RP2005859	14954	14955
NT2RP2006023	C-NT2RP2006023	14956	14957

(970)

特開 2002-191363

1937

1938

NT2RP2006334	C-NT2RP2006334	14958	14959
NT2RP2006441	C-NT2RP2006441	14960	
NT2RP3000002	C-NT2RP3000002	14961	14962
NT2RP3000050	C-NT2RP3000050	14963	14964
NT2RP3000055	C-NT2RP3000055	14965	14966
NT2RP3000068	C-NT2RP3000068	14967	14968
NT2RP3000080	C-NT2RP3000080	14969	14970
NT2RP3000085	C-NT2RP3000085	14971	14972
NT2RP3000092	C-NT2RP3000092	14973	14974
NT2RP3000109	C-NT2RP3000109	14975	14976
NT2RP3000134	C-NT2RP3000134	14977	14978
NT2RP3000149	C-NT2RP3000149	14979	14980
NT2RP3000197	C-NT2RP3000197	14981	
NT2RP3000207	C-NT2RP3000207	14982	14983
NT2RP3000233	C-NT2RP3000233	14984	14985
NT2RP3000235	C-NT2RP3000235	14986	
NT2RP3000247	C-NT2RP3000247	14987	
NT2RP3000267	C-NT2RP3000267	14988	14989
NT2RP3000299	C-NT2RP3000299	14990	14991
NT2RP3000324	C-NT2RP3000324	14992	
NT2RP3000341	C-NT2RP3000341	14993	
NT2RP3000393	C-NT2RP3000393	14994	14995
NT2RP3000441	C-NT2RP3000441	14996	14997
NT2RP3000449	C-NT2RP3000449	14998	14999
NT2RP3000451	C-NT2RP3000451	15000	15001
NT2RP3000456	C-NT2RP3000456	15002	15003
NT2RP3000542	C-NT2RP3000542	15004	
NT2RP3000561	C-NT2RP3000561	15005	
NT2RP3000562	C-NT2RP3000562	15006	
NT2RP3000578	C-NT2RP3000578	15007	15008
NT2RP3000590	C-NT2RP3000590	15009	15010
NT2RP3000592	C-NT2RP3000592	15011	
NT2RP3000622	C-NT2RP3000622	15012	
NT2RP3000624	C-NT2RP3000624	15013	15014
NT2RP3000685	C-NT2RP3000685	15015	15016
NT2RP3000736	C-NT2RP3000736	15017	15018
NT2RP3000742	C-NT2RP3000742	15019	15020
NT2RP3000753	C-NT2RP3000753	15021	15022
NT2RP3000826	C-NT2RP3000826	15023	15024
NT2RP3000865	C-NT2RP3000865	15025	
NT2RP3000875	C-NT2RP3000875	15026	15027
NT2RP3001007	C-NT2RP3001007	15028	
NT2RP3001055	C-NT2RP3001055	15029	15030
NT2RP3001111	C-NT2RP3001111	15031	15032
NT2RP3001120	C-NT2RP3001120	15033	15034
NT2RP3001126	C-NT2RP3001126	15035	15036
NT2RP3001150	C-NT2RP3001150	15037	15038
NT2RP3001232	C-NT2RP3001232	15039	
NT2RP3001268	C-NT2RP3001268	15040	15041
NT2RP3001272	C-NT2RP3001272	15042	15043

(971)

特開 2002-191363

1939

1940

NT2RP3001274	C-NT2RP3001274	15044	15045
NT2RP3001281	C-NT2RP3001281	15046	15047
NT2RP3001297	C-NT2RP3001297	15048	15049
NT2RP3001318	C-NT2RP3001318	15050	
NT2RP3001338	C-NT2RP3001338	15051	15052
NT2RP3001355	C-NT2RP3001355	15053	15054
NT2RP3001374	C-NT2RP3001374	15055	15056
NT2RP3001428	C-NT2RP3001428	15057	15058
NT2RP3001432	C-NT2RP3001432	15059	
NT2RP3001447	C-NT2RP3001447	15060	15061
NT2RP3001449	C-NT2RP3001449	15062	15063
NT2RP3001453	C-NT2RP3001453	15064	15065
NT2RP3001459	C-NT2RP3001459	15066	
NT2RP3001527	C-NT2RP3001527	15067	15068
NT2RP3001538	C-NT2RP3001538	15069	15070
NT2RP3001580	C-NT2RP3001580	15071	15072
NT2RP3001587	C-NT2RP3001587	15073	15074
NT2RP3001589	C-NT2RP3001589	15075	15076
NT2RP3001607	C-NT2RP3001607	15077	
NT2RP3001608	C-NT2RP3001608	15078	15079
NT2RP3001671	C-NT2RP3001671	15080	15081
NT2RP3001672	C-NT2RP3001672	15082	15083
NT2RP3001678	C-NT2RP3001678	15084	15085
NT2RP3001688	C-NT2RP3001688	15086	15087
NT2RP3001690	C-NT2RP3001690	15088	15089
NT2RP3001698	C-NT2RP3001698	15090	
NT2RP3001708	C-NT2RP3001708	15091	15092
NT2RP3001716	C-NT2RP3001716	15093	15094
NT2RP3001752	C-NT2RP3001752	15095	
NT2RP3001792	C-NT2RP3001792	15096	15097
NT2RP3001844	C-NT2RP3001844	15098	
NT2RP3001854	C-NT2RP3001854	15099	15100
NT2RP3001855	C-NT2RP3001855	15101	15102
NT2RP3001898	C-NT2RP3001898	15103	15104
NT2RP3001931	C-NT2RP3001931	15105	15106
NT2RP3001969	C-NT2RP3001969	15107	15108
NT2RP3002002	C-NT2RP3002002	15109	15110
NT2RP3002004	C-NT2RP3002004	15111	15112
NT2RP3002007	C-NT2RP3002007	15113	15114
NT2RP3002014	C-NT2RP3002014	15115	15116
NT2RP3002045	C-NT2RP3002045	15117	15118
NT2RP3002056	C-NT2RP3002056	15119	15120
NT2RP3002062	C-NT2RP3002062	15121	15122
NT2RP3002081	C-NT2RP3002081	15123	15124
NT2RP3002097	C-NT2RP3002097	15125	15126
NT2RP3002108	C-NT2RP3002108	15127	15128
NT2RP3002142	C-NT2RP3002142	15129	
NT2RP3002146	C-NT2RP3002146	15130	15131
NT2RP3002151	C-NT2RP3002151	15132	15133
NT2RP3002165	C-NT2RP3002165	15134	15135

(972)

特開 2002-191363

1941

1942

NT2RP3002166	C-NT2RP3002166	15136	15137
NT2RP3002181	C-NT2RP3002181	15138	15139
NT2RP3002244	C-NT2RP3002244	15140	15141
NT2RP3002248	C-NT2RP3002248	15142	15143
NT2RP3002273	C-NT2RP3002273	15144	15145
NT2RP3002276	C-NT2RP3002276	15146	
NT2RP3002304	C-NT2RP3002304	15147	
NT2RP3002501	C-NT2RP3002501	15148	15149
NT2RP3002529	C-NT2RP3002529	15150	15151
NT2RP3002566	C-NT2RP3002566	15152	15153
NT2RP3002587	C-NT2RP3002587	15154	
NT2RP3002590	C-NT2RP3002590	15155	15156
NT2RP3002631	C-NT2RP3002631	15157	15158
NT2RP3002650	C-NT2RP3002650	15159	15160
NT2RP3002663	C-NT2RP3002663	15161	15162
NT2RP3002671	C-NT2RP3002671	15163	15164
NT2RP3002763	C-NT2RP3002763	15165	15166
NT2RP3002861	C-NT2RP3002861	15167	
NT2RP3002911	C-NT2RP3002911	15168	
NT2RP3002948	C-NT2RP3002948	15169	15170
NT2RP3002953	C-NT2RP3002953	15171	15172
NT2RP3002988	C-NT2RP3002988	15173	15174
NT2RP3003008	C-NT2RP3003008	15175	15176
NT2RP3003101	C-NT2RP3003101	15177	15178
NT2RP3003204	C-NT2RP3003204	15179	15180
NT2RP3003278	C-NT2RP3003278	15181	15182
NT2RP3003282	C-NT2RP3003282	15183	15184
NT2RP3003290	C-NT2RP3003290	15185	15186
NT2RP3003302	C-NT2RP3003302	15187	15188
NT2RP3003313	C-NT2RP3003313	15189	15190
NT2RP3003327	C-NT2RP3003327	15191	
NT2RP3003344	C-NT2RP3003344	15192	15193
NT2RP3003353	C-NT2RP3003353	15194	15195
NT2RP3003377	C-NT2RP3003377	15196	15197
NT2RP3003385	C-NT2RP3003385	15198	15199
NT2RP3003433	C-NT2RP3003433	15200	
NT2RP3003490	C-NT2RP3003490	15201	15202
NT2RP3003491	C-NT2RP3003491	15203	15204
NT2RP3004206	C-NT2RP3004206	15205	15206
NT2RP3004207	C-NT2RP3004207	15207	15208
NT2RP3004209	C-NT2RP3004209	15209	15210
NT2RP3004242	C-NT2RP3004242	15211	15212
NT2RP3004246	C-NT2RP3004246	15213	15214
NT2RP3004258	C-NT2RP3004258	15215	15216
NT2RP3004262	C-NT2RP3004262	15217	15218
NT2RP3004341	C-NT2RP3004341	15219	
NT2RP3004378	C-NT2RP3004378	15220	15221
NT2RP3004424	C-NT2RP3004424	15222	15223
NT2RP3004428	C-NT2RP3004428	15224	15225
NT2RP3004451	C-NT2RP3004451	15226	15227

(973)

特開 2002-191363

1943

1944

NT2RP3004454	C-NT2RP3004454	15228	15229
NT2RP3004472	C-NT2RP3004472	15230	15231
NT2RP3004498	C-NT2RP3004498	15232	15233
NT2RP3004504	C-NT2RP3004504	15234	15235
NT2RP3004507	C-NT2RP3004507	15236	15237
NT2RP3004534	C-NT2RP3004534	15238	15239
NT2RP4000528	C-NT2RP4000528	15240	15241
NT2RP4000907	C-NT2RP4000907	15242	15243
NT2RP4001029	C-NT2RP4001029	15244	15245
NT2RP4001336	C-NT2RP4001336	15246	15247
NT2RP4001389	C-NT2RP4001389	15248	15249
NT2RP4001442	C-NT2RP4001442	15250	15251
NT2RP4001529	C-NT2RP4001529	15252	15253
NT2RP4001656	C-NT2RP4001656	15254	15255
OVARC1000106	C-OVARC1000106	15256	15257
OVARC1000198	C-OVARC1000198	15258	
OVARC1000682	C-OVARC1000682	15259	15260
OVARC1000703	C-OVARC1000703	15261	15262
OVARC1000722	C-OVARC1000722	15263	15264
OVARC1000730	C-OVARC1000730	15265	15266
OVARC1000746	C-OVARC1000746	15267	15268
OVARC1000781	C-OVARC1000781	15269	15270
OVARC1000787	C-OVARC1000787	15271	
OVARC1000834	C-OVARC1000834	15272	15273
OVARC1000846	C-OVARC1000846	15274	15275
OVARC1000850	C-OVARC1000850	15276	15277
OVARC1000862	C-OVARC1000862	15278	15279
OVARC1000876	C-OVARC1000876	15280	15281
OVARC1000883	C-OVARC1000883	15282	
OVARC1000886	C-OVARC1000886	15283	15284
OVARC1000912	C-OVARC1000912	15285	15286
OVARC1000915	C-OVARC1000915	15287	15288
OVARC1000924	C-OVARC1000924	15289	15290
OVARC1000964	C-OVARC1000964	15291	
OVARC1000984	C-OVARC1000984	15292	15293
OVARC1001004	C-OVARC1001004	15294	
OVARC1001010	C-OVARC1001010	15295	15296
OVARC1001011	C-OVARC1001011	15297	
OVARC1001032	C-OVARC1001032	15298	15299
OVARC1001044	C-OVARC1001044	15300	
OVARC1001055	C-OVARC1001055	15301	
OVARC1001068	C-OVARC1001068	15302	15303
OVARC1001074	C-OVARC1001074	15304	
OVARC1001092	C-OVARC1001092	15305	15306
OVARC1001107	C-OVARC1001107	15307	15308
OVARC1001154	C-OVARC1001154	15309	15310
OVARC1001161	C-OVARC1001161	15311	15312
OVARC1001167	C-OVARC1001167	15313	15314
OVARC1001170	C-OVARC1001170	15315	
OVARC1001171	C-OVARC1001171	15316	15317

(974)

特開 2002-191363

1945

1946

OVARC1001173	C-OVARC1001173	15318	
OVARC1001176	C-OVARC1001176	15319	15320
OVARC1001180	C-OVARC1001180	15321	15322
OVARC1001188	C-OVARC1001188	15323	15324
OVARC1001232	C-OVARC1001232	15325	15326
OVARC1001270	C-OVARC1001270	15327	15328
OVARC1001271	C-OVARC1001271	15329	15330
OVARC1001306	C-OVARC1001306	15331	15332
OVARC1001344	C-OVARC1001344	15333	
OVARC1001369	C-OVARC1001369	15334	15335
OVARC1001372	C-OVARC1001372	15336	15337
OVARC1001391	C-OVARC1001391	15338	15339
OVARC1001399	C-OVARC1001399	15340	15341
OVARC1001417	C-OVARC1001417	15342	15343
OVARC1001419	C-OVARC1001419	15344	15345
OVARC1001436	C-OVARC1001436	15346	15347
OVARC1001453	C-OVARC1001453	15348	15349
OVARC1001476	C-OVARC1001476	15350	15351
OVARC1001480	C-OVARC1001480	15352	15353
OVARC1001489	C-OVARC1001489	15354	
OVARC1001506	C-OVARC1001506	15355	15356
OVARC1001525	C-OVARC1001525	15357	15358
OVARC1001555	C-OVARC1001555	15359	15360
OVARC1001577	C-OVARC1001577	15361	15362
OVARC1001600	C-OVARC1001600	15363	
OVARC1001610	C-OVARC1001610	15364	15365
OVARC1001702	C-OVARC1001702	15366	15367
OVARC1001703	C-OVARC1001703	15368	15369
OVARC1001711	C-OVARC1001711	15370	15371
OVARC1001713	C-OVARC1001713	15372	15373
OVARC1001726	C-OVARC1001726	15374	15375
OVARC1001731	C-OVARC1001731	15376	15377
OVARC1001745	C-OVARC1001745	15378	15379
OVARC1001762	C-OVARC1001762	15380	15381
OVARC1001766	C-OVARC1001766	15382	15383
OVARC1001767	C-OVARC1001767	15384	15385
OVARC1001768	C-OVARC1001768	15386	15387
OVARC1001791	C-OVARC1001791	15388	15389
OVARC1001795	C-OVARC1001795	15390	
OVARC1001802	C-OVARC1001802	15391	15392
OVARC1001809	C-OVARC1001809	15393	15394
OVARC1001828	C-OVARC1001828	15395	
OVARC1001846	C-OVARC1001846	15396	
OVARC1001861	C-OVARC1001861	15397	15398
OVARC1001879	C-OVARC1001879	15399	15400
OVARC1001880	C-OVARC1001880	15401	15402
OVARC1001883	C-OVARC1001883	15403	
OVARC1001916	C-OVARC1001916	15404	15405
OVARC1001928	C-OVARC1001928	15406	
OVARC1001942	C-OVARC1001942	15407	15408

(975)

特開 2002-191363

1947

1948

OVARC1001943	C-OVARC1001943	15409	15410
OVARC1001950	C-OVARC1001950	15411	
OVARC1001987	C-OVARC1001987	15412	15413
OVARC1002050	C-OVARC1002050	15414	15415
OVARC1002082	C-OVARC1002082	15416	
OVARC1002107	C-OVARC1002107	15417	15418
OVARC1002127	C-OVARC1002127	15419	15420
OVARC1002138	C-OVARC1002138	15421	15422
OVARC1002156	C-OVARC1002156	15423	15424
OVARC1002158	C-OVARC1002158	15425	15426
PLACE1000004	C-PLACE1000004	15427	15428
PLACE1000040	C-PLACE1000040	15429	15430
PLACE1000048	C-PLACE1000048	15431	15432
PLACE1000050	C-PLACE1000050	15433	15434
PLACE1000061	C-PLACE1000061	15435	
PLACE1000081	C-PLACE1000081	15436	15437
PLACE1000094	C-PLACE1000094	15438	
PLACE1000133	C-PLACE1000133	15439	15440
PLACE1000214	C-PLACE1000214	15441	
PLACE1000236	C-PLACE1000236	15442	
PLACE1000246	C-PLACE1000246	15443	15444
PLACE1000292	C-PLACE1000292	15445	
PLACE1000308	C-PLACE1000308	15446	
PLACE1000332	C-PLACE1000332	15447	
PLACE1000453	C-PLACE1000453	15448	
PLACE1000583	C-PLACE1000583	15449	
PLACE1000599	C-PLACE1000599	15450	
PLACE1000610	C-PLACE1000610	15451	15452
PLACE1000653	C-PLACE1000653	15453	15454
PLACE1000656	C-PLACE1000656	15455	15456
PLACE1000706	C-PLACE1000706	15457	15458
PLACE1000712	C-PLACE1000712	15459	15460
PLACE1000749	C-PLACE1000749	15461	15462
PLACE1000769	C-PLACE1000769	15463	15464
PLACE1000786	C-PLACE1000786	15465	15466
PLACE1000849	C-PLACE1000849	15467	15468
PLACE1000856	C-PLACE1000856	15469	15470
PLACE1000931	C-PLACE1000931	15471	15472
PLACE1000987	C-PLACE1000987	15473	15474
PLACE1001010	C-PLACE1001010	15475	15476
PLACE1001015	C-PLACE1001015	15477	
PLACE1001024	C-PLACE1001024	15478	15479
PLACE1001062	C-PLACE1001062	15480	15481
PLACE1001104	C-PLACE1001104	15482	15483
PLACE1001168	C-PLACE1001168	15484	
PLACE1001171	C-PLACE1001171	15485	15486
PLACE1001185	C-PLACE1001185	15487	15488
PLACE1001238	C-PLACE1001238	15489	15490
PLACE1001280	C-PLACE1001280	15491	
PLACE1001294	C-PLACE1001294	15492	15493

(976)

特開 2002-191363

1949

1950

PLACE1001304	C-PLACE1001304	15494	15495
PLACE1001311	C-PLACE1001311	15496	
PLACE1001323	C-PLACE1001323	15497	
PLACE1001351	C-PLACE1001351	15498	15499
PLACE1001414	C-PLACE1001414	15500	
PLACE1001440	C-PLACE1001440	15501	
PLACE1001456	C-PLACE1001456	15502	
PLACE1001517	C-PLACE1001517	15503	15504
PLACE1001602	C-PLACE1001602	15505	15506
PLACE1001632	C-PLACE1001632	15507	15508
PLACE1001634	C-PLACE1001634	15509	
PLACE1001640	C-PLACE1001640	15510	
PLACE1001672	C-PLACE1001672	15511	15512
PLACE1001705	C-PLACE1001705	15513	
PLACE1001716	C-PLACE1001716	15514	15515
PLACE1001720	C-PLACE1001720	15516	15517
PLACE1001745	C-PLACE1001745	15518	15519
PLACE1001748	C-PLACE1001748	15520	15521
PLACE1001771	C-PLACE1001771	15522	15523
PLACE1001799	C-PLACE1001799	15524	
PLACE1001845	C-PLACE1001845	15525	15526
PLACE1001897	C-PLACE1001897	15527	
PLACE1002090	C-PLACE1002090	15528	
PLACE1002157	C-PLACE1002157	15529	15530
PLACE1002171	C-PLACE1002171	15531	15532
PLACE1002227	C-PLACE1002227	15533	
PLACE1002259	C-PLACE1002259	15534	
PLACE1002319	C-PLACE1002319	15535	15536
PLACE1002395	C-PLACE1002395	15537	15538
PLACE1002477	C-PLACE1002477	15539	
PLACE1002493	C-PLACE1002493	15540	15541
PLACE1002500	C-PLACE1002500	15542	15543
PLACE1002514	C-PLACE1002514	15544	
PLACE1002532	C-PLACE1002532	15545	15546
PLACE1002537	C-PLACE1002537	15547	
PLACE1002571	C-PLACE1002571	15548	15549
PLACE1002583	C-PLACE1002583	15550	
PLACE1002598	C-PLACE1002598	15551	
PLACE1002625	C-PLACE1002625	15552	15553
PLACE1002655	C-PLACE1002655	15554	15555
PLACE1002768	C-PLACE1002768	15556	
PLACE1002782	C-PLACE1002782	15557	
PLACE1002816	C-PLACE1002816	15558	15559
PLACE1002853	C-PLACE1002853	15560	
PLACE1002908	C-PLACE1002908	15561	15562
PLACE1002962	C-PLACE1002962	15563	
PLACE1002968	C-PLACE1002968	15564	
PLACE1002991	C-PLACE1002991	15565	15566
PLACE1003025	C-PLACE1003025	15567	15568
PLACE1003027	C-PLACE1003027	15569	15570

(977)

特開 2002-191363

1951

1952

PLACE1003044	C-PLACE1003044	15571	15572
PLACE1003176	C-PLACE1003176	15573	15574
PLACE1003238	C-PLACE1003238	15575	15576
PLACE1003256	C-PLACE1003256	15577	
PLACE1003258	C-PLACE1003258	15578	15579
PLACE1003343	C-PLACE1003343	15580	
PLACE1003361	C-PLACE1003361	15581	
PLACE1003366	C-PLACE1003366	15582	15583
PLACE1003373	C-PLACE1003373	15584	
PLACE1003375	C-PLACE1003375	15585	15586
PLACE1003394	C-PLACE1003394	15587	15588
PLACE1003420	C-PLACE1003420	15589	15590
PLACE1003454	C-PLACE1003454	15591	
PLACE1003478	C-PLACE1003478	15592	
PLACE1003516	C-PLACE1003516	15593	
PLACE1003519	C-PLACE1003519	15594	15595
PLACE1003521	C-PLACE1003521	15596	15597
PLACE1003528	C-PLACE1003528	15598	
PLACE1003537	C-PLACE1003537	15599	15600
PLACE1003566	C-PLACE1003566	15601	
PLACE1003584	C-PLACE1003584	15602	
PLACE1003593	C-PLACE1003593	15603	
PLACE1003605	C-PLACE1003605	15604	15605
PLACE1003618	C-PLACE1003618	15606	
PLACE1003638	C-PLACE1003638	15607	
PLACE1003738	C-PLACE1003738	15608	15609
PLACE1003760	C-PLACE1003760	15610	
PLACE1003768	C-PLACE1003768	15611	15612
PLACE1003795	C-PLACE1003795	15613	15614
PLACE1003886	C-PLACE1003886	15615	15616
PLACE1003888	C-PLACE1003888	15617	15618
PLACE1003903	C-PLACE1003903	15619	15620
PLACE1003915	C-PLACE1003915	15621	15622
PLACE1004118	C-PLACE1004118	15623	15624
PLACE1004256	C-PLACE1004256	15625	15626
PLACE1004274	C-PLACE1004274	15627	15628
PLACE1004284	C-PLACE1004284	15629	
PLACE1005331	C-PLACE1005331	15630	15631
PLACE1005739	C-PLACE1005739	15632	15633
PLACE1005828	C-PLACE1005828	15634	15635
PLACE1005876	C-PLACE1005876	15636	15637
PLACE1005890	C-PLACE1005890	15638	15639
PLACE1006157	C-PLACE1006157	15640	15641
PLACE1007053	C-PLACE1007053	15642	15643
PLACE1007068	C-PLACE1007068	15644	15645
PLACE1008368	C-PLACE1008368	15646	15647
PLACE1009921	C-PLACE1009921	15648	
PLACE1010401	C-PLACE1010401	15649	
PLACE1010856	C-PLACE1010856	15650	
PLACE1010857	C-PLACE1010857	15651	15652

(978)

特開 2002-191363

1953

1954

PLACE1010917	C-PLACE1010917	15653	15654
PLACE1010925	C-PLACE1010925	15655	
PLACE1010926	C-PLACE1010926	15656	15657
PLACE1010942	C-PLACE1010942	15658	15659
PLACE1010944	C-PLACE1010944	15660	
PLACE1010954	C-PLACE1010954	15661	15662
PLACE1010960	C-PLACE1010960	15663	15664
PLACE1011026	C-PLACE1011026	15665	15666
PLACE1011046	C-PLACE1011046	15667	15668
PLACE1011054	C-PLACE1011054	15669	
PLACE1011057	C-PLACE1011057	15670	
PLACE1011109	C-PLACE1011109	15671	
PLACE1011114	C-PLACE1011114	15672	15673
PLACE1011133	C-PLACE1011133	15674	
PLACE1011143	C-PLACE1011143	15675	
PLACE1011165	C-PLACE1011165	15676	
PLACE1011185	C-PLACE1011185	15677	15678
PLACE1011219	C-PLACE1011219	15679	15680
PLACE1011221	C-PLACE1011221	15681	15682
PLACE1011263	C-PLACE1011263	15683	15684
PLACE1011325	C-PLACE1011325	15685	15686
PLACE1011332	C-PLACE1011332	15687	15688
PLACE1011340	C-PLACE1011340	15689	
PLACE1011399	C-PLACE1011399	15690	15691
PLACE1011433	C-PLACE1011433	15692	15693
PLACE1011452	C-PLACE1011452	15694	
PLACE1011465	C-PLACE1011465	15695	15696
PLACE1011472	C-PLACE1011472	15697	15698
PLACE1011477	C-PLACE1011477	15699	15700
PLACE1011492	C-PLACE1011492	15701	15702
PLACE1011520	C-PLACE1011520	15703	
PLACE1011563	C-PLACE1011563	15704	15705
PLACE1011567	C-PLACE1011567	15706	
PLACE1011576	C-PLACE1011576	15707	15708
PLACE1011586	C-PLACE1011586	15709	15710
PLACE1011643	C-PLACE1011643	15711	15712
PLACE1011649	C-PLACE1011649	15713	15714
PLACE1011664	C-PLACE1011664	15715	15716
PLACE1011682	C-PLACE1011682	15717	
PLACE1011719	C-PLACE1011719	15718	
PLACE1011729	C-PLACE1011729	15719	
PLACE1011858	C-PLACE1011858	15720	15721
PLACE1011874	C-PLACE1011874	15722	
PLACE1011875	C-PLACE1011875	15723	15724
PLACE1011923	C-PLACE1011923	15725	15726
PLACE1011982	C-PLACE1011982	15727	
PLACE2000014	C-PLACE2000014	15728	
PLACE2000015	C-PLACE2000015	15729	15730
PLACE2000017	C-PLACE2000017	15731	
PLACE2000021	C-PLACE2000021	15732	15733

(979)

特開 2002-191363

1955

1956

PLACE2000047	C-PLACE2000047	15734	
PLACE2000062	C-PLACE2000062	15735	15736
PLACE2000100	C-PLACE2000100	15737	
PLACE2000111	C-PLACE2000111	15738	15739
PLACE2000172	C-PLACE2000172	15740	15741
PLACE2000187	C-PLACE2000187	15742	15743
PLACE2000216	C-PLACE2000216	15744	15745
PLACE2000246	C-PLACE2000246	15746	15747
PLACE2000317	C-PLACE2000317	15748	
PLACE2000341	C-PLACE2000341	15749	15750
PLACE2000366	C-PLACE2000366	15751	
PLACE2000373	C-PLACE2000373	15752	15753
PLACE2000394	C-PLACE2000394	15754	15755
PLACE2000398	C-PLACE2000398	15756	15757
PLACE2000411	C-PLACE2000411	15758	15759
PLACE2000425	C-PLACE2000425	15760	15761
PLACE2000427	C-PLACE2000427	15762	15763
PLACE2000433	C-PLACE2000433	15764	
PLACE2000438	C-PLACE2000438	15765	15766
PLACE2000458	C-PLACE2000458	15767	15768
PLACE2000477	C-PLACE2000477	15769	15770
PLACE3000009	C-PLACE3000009	15771	15772
PLACE3000020	C-PLACE3000020	15773	15774
PLACE3000103	C-PLACE3000103	15775	15776
PLACE3000142	C-PLACE3000142	15777	
PLACE3000145	C-PLACE3000145	15778	15779
PLACE3000156	C-PLACE3000156	15780	
PLACE3000157	C-PLACE3000157	15781	15782
PLACE3000197	C-PLACE3000197	15783	15784
PLACE3000208	C-PLACE3000208	15785	15786
PLACE3000226	C-PLACE3000226	15787	15788
PLACE3000242	C-PLACE3000242	15789	15790
PLACE3000363	C-PLACE3000363	15791	15792
PLACE3000405	C-PLACE3000405	15793	15794
PLACE3000416	C-PLACE3000416	15795	15796
PLACE3000477	C-PLACE3000477	15797	15798
PLACE4000106	C-PLACE4000106	15799	15800
PLACE4000323	C-PLACE4000323	15801	
PLACE4000326	C-PLACE4000326	15802	15803
PLACE4000369	C-PLACE4000369	15804	15805
PLACE4000445	C-PLACE4000445	15806	15807
PLACE4000558	C-PLACE4000558	15808	15809
PLACE4000581	C-PLACE4000581	15810	15811
PLACE4000593	C-PLACE4000593	15812	15813
PLACE4000612	C-PLACE4000612	15814	15815
PLACE4000670	C-PLACE4000670	15816	15817
THYR01000026	C-THYR01000026	15818	
THYR01000085	C-THYR01000085	15819	
THYR01000107	C-THYR01000107	15820	
THYR01000111	C-THYR01000111	15821	15822

(980)

特開 2002-191363

1957

1958

THYRO1000132	C-THYRO1000132	15823	15824
THYRO1000156	C-THYRO1000156	15825	15826
THYRO1000173	C-THYRO1000173	15827	15828
THYRO1000186	C-THYRO1000186	15829	
THYRO1000187	C-THYRO1000187	15830	
THYRO1000241	C-THYRO1000241	15831	
THYRO1000279	C-THYRO1000279	15832	
THYRO1000327	C-THYRO1000327	15833	15834
THYRO1000452	C-THYRO1000452	15835	
THYRO1000471	C-THYRO1000471	15836	
THYRO1000484	C-THYRO1000484	15837	15838
THYRO1000502	C-THYRO1000502	15839	15840
THYRO1000505	C-THYRO1000505	15841	15842
THYRO1000585	C-THYRO1000585	15843	15844
THYRO1000596	C-THYRO1000596	15845	
THYRO1000662	C-THYRO1000662	15846	15847
THYRO1000666	C-THYRO1000666	15848	15849
THYRO1000715	C-THYRO1000715	15850	15851
THYRO1000734	C-THYRO1000734	15852	
THYRO1000748	C-THYRO1000748	15853	15854
THYRO1000756	C-THYRO1000756	15855	15856
THYRO1000777	C-THYRO1000777	15857	15858
THYRO1000783	C-THYRO1000783	15859	15860
THYRO1000787	C-THYRO1000787	15861	15862
THYRO1000793	C-THYRO1000793	15863	15864
THYRO1000796	C-THYRO1000796	15865	
THYRO1000843	C-THYRO1000843	15866	
THYRO1000852	C-THYRO1000852	15867	15868
THYRO1000865	C-THYRO1000865	15869	
THYRO1000895	C-THYRO1000895	15870	15871
THYRO1000926	C-THYRO1000926	15872	15873
THYRO1000951	C-THYRO1000951	15874	15875
THYRO1000952	C-THYRO1000952	15876	15877
THYRO1000983	C-THYRO1000983	15878	15879
THYRO1001003	C-THYRO1001003	15880	
THYRO1001031	C-THYRO1001031	15881	
THYRO1001062	C-THYRO1001062	15882	
THYRO1001100	C-THYRO1001100	15883	15884
THYRO1001133	C-THYRO1001133	15885	15886
THYRO1001134	C-THYRO1001134	15887	15888
THYRO1001173	C-THYRO1001173	15889	15890
THYRO1001213	C-THYRO1001213	15891	
THYRO1001321	C-THYRO1001321	15892	
THYRO1001322	C-THYRO1001322	15893	15894
THYRO1001365	C-THYRO1001365	15895	15896
THYRO1001401	C-THYRO1001401	15897	15898
THYRO1001411	C-THYRO1001411	15899	
THYRO1001434	C-THYRO1001434	15900	
THYRO1001534	C-THYRO1001534	15901	15902
THYRO1001541	C-THYRO1001541	15903	15904

(981)

特開 2002-191363

1959

1960

THYRO1001559	C-THYRO1001559	15905	
THYRO1001570	C-THYRO1001570	15906	15907
THYRO1001595	C-THYRO1001595	15908	
THYRO1001605	C-THYRO1001605	15909	
THYRO1001617	C-THYRO1001617	15910	15911
THYRO1001656	C-THYRO1001656	15912	15913
THYRO1001671	C-THYRO1001671	15914	15915
THYRO1001673	C-THYRO1001673	15916	
THYRO1001703	C-THYRO1001703	15917	15918
THYRO1001706	C-THYRO1001706	15919	
THYRO1001738	C-THYRO1001738	15920	15921
THYRO1001745	C-THYRO1001745	15922	15923
THYRO1001793	C-THYRO1001793	15924	15925
THYRO1001809	C-THYRO1001809	15926	15927
THYRO1001895	C-THYRO1001895	15928	
THYRO1001907	C-THYRO1001907	15929	
VESEN1000122	C-VESEN1000122	15930	15931
Y79AA1000037	C-Y79AA1000037	15932	15933
Y79AA1000059	C-Y79AA1000059	15934	15935
Y79AA1000065	C-Y79AA1000065	15936	15937
Y79AA1000131	C-Y79AA1000131	15938	15939
Y79AA1000181	C-Y79AA1000181	15940	15941
Y79AA1000202	C-Y79AA1000202	15942	15943
Y79AA1000214	C-Y79AA1000214	15944	15945
Y79AA1000230	C-Y79AA1000230	15946	15947
Y79AA1000258	C-Y79AA1000258	15948	15949
Y79AA1000268	C-Y79AA1000268	15950	15951
Y79AA1000313	C-Y79AA1000313	15952	15953
Y79AA1000328	C-Y79AA1000328	15954	15955
Y79AA1000355	C-Y79AA1000355	15956	15957
Y79AA1000368	C-Y79AA1000368	15958	15959
Y79AA1000420	C-Y79AA1000420	15960	15961
Y79AA1000469	C-Y79AA1000469	15962	15963
Y79AA1000480	C-Y79AA1000480	15964	15965
Y79AA1000540	C-Y79AA1000540	15966	15967
Y79AA1000560	C-Y79AA1000560	15968	15969
Y79AA1000574	C-Y79AA1000574	15970	15971
Y79AA1000627	C-Y79AA1000627	15972	15973
Y79AA1000705	C-Y79AA1000705	15974	15975
Y79AA1000734	C-Y79AA1000734	15976	15977
Y79AA1000748	C-Y79AA1000748	15978	15979
Y79AA1000752	C-Y79AA1000752	15980	15981
Y79AA1000774	C-Y79AA1000774	15982	15983
Y79AA1000782	C-Y79AA1000782	15984	15985
Y79AA1000784	C-Y79AA1000784	15986	15987
Y79AA1000794	C-Y79AA1000794	15988	15989
Y79AA1000800	C-Y79AA1000800	15990	15991
Y79AA1000805	C-Y79AA1000805	15992	
Y79AA1000824	C-Y79AA1000824	15993	15994
Y79AA1000833	C-Y79AA1000833	15995	15996

(982)

特開 2002-191363

1961

1962

Y79AA1000850	C-Y79AA1000850	15997	15998
Y79AA1000962	C-Y79AA1000962	15999	16000
Y79AA1000968	C-Y79AA1000968	16001	16002
Y79AA1000976	C-Y79AA1000976	16003	16004
Y79AA1001023	C-Y79AA1001023	16005	16006
Y79AA1001041	C-Y79AA1001041	16007	16008
Y79AA1001048	C-Y79AA1001048	16009	16010
Y79AA1001077	C-Y79AA1001077	16011	16012
Y79AA1001078	C-Y79AA1001078	16013	16014
Y79AA1001145	C-Y79AA1001145	16015	16016
Y79AA1001177	C-Y79AA1001177	16017	16018
Y79AA1001185	C-Y79AA1001185	16019	16020
Y79AA1001211	C-Y79AA1001211	16021	16022
Y79AA1001228	C-Y79AA1001228	16023	16024
Y79AA1001233	C-Y79AA1001233	16025	16026
Y79AA1001236	C-Y79AA1001236	16027	16028
Y79AA1001281	C-Y79AA1001281	16029	16030
Y79AA1001312	C-Y79AA1001312	16031	16032
Y79AA1001323	C-Y79AA1001323	16033	16034
Y79AA1001391	C-Y79AA1001391	16035	16036
Y79AA1001394	C-Y79AA1001394	16037	16038
Y79AA1001402	C-Y79AA1001402	16039	16040
Y79AA1001493	C-Y79AA1001493	16041	16042
Y79AA1001533	C-Y79AA1001533	16043	16044
Y79AA1001541	C-Y79AA1001541	16045	16046
Y79AA1001548	C-Y79AA1001548	16047	
Y79AA1001555	C-Y79AA1001555	16048	
Y79AA1001581	C-Y79AA1001581	16049	16050
Y79AA1001585	C-Y79AA1001585	16051	
Y79AA1001603	C-Y79AA1001603	16052	16053
Y79AA1001613	C-Y79AA1001613	16054	16055
Y79AA1001665	C-Y79AA1001665	16056	
Y79AA1001679	C-Y79AA1001679	16057	16058
Y79AA1001696	C-Y79AA1001696	16059	16060
Y79AA1001705	C-Y79AA1001705	16061	16062
Y79AA1001711	C-Y79AA1001711	16063	16064
Y79AA1001781	C-Y79AA1001781	16065	16066
Y79AA1001805	C-Y79AA1001805	16067	16068
Y79AA1001827	C-Y79AA1001827	16069	16070
Y79AA1001846	C-Y79AA1001846	16071	16072
Y79AA1001923	C-Y79AA1001923	16073	16074
Y79AA1001963	C-Y79AA1001963	16075	16076
Y79AA1002027	C-Y79AA1002027	16077	16078
Y79AA1002083	C-Y79AA1002083	16079	16080
Y79AA1002089	C-Y79AA1002089	16081	16082
Y79AA1002115	C-Y79AA1002115	16083	16084
Y79AA1002125	C-Y79AA1002125	16085	16086
Y79AA1002204	C-Y79AA1002204	16087	16088
Y79AA1002208	C-Y79AA1002208	16089	16090
Y79AA1002209	C-Y79AA1002209	16091	16092

(983)

特開 2002-191363

1963

1964

Y79AA1002229	C-Y79AA1002229	16093	16094
Y79AA1002246	C-Y79AA1002246	16095	16096
Y79AA1002298	C-Y79AA1002298	16097	16098
Y79AA1002307	C-Y79AA1002307	16099	16100
Y79AA1002311	C-Y79AA1002311	16101	16102
Y79AA1002351	C-Y79AA1002351	16103	16104
Y79AA1002407	C-Y79AA1002407	16105	16106
Y79AA1002433	C-Y79AA1002433	16107	16108
Y79AA1002472	C-Y79AA1002472	16109	16110
HEMBA1000042	C-HEMBA1000042	16219	
HEMBA1000141	C-HEMBA1000141	16220	16221
HEMBA1000150	C-HEMBA1000150	16222	16223
HEMBA1000213	C-HEMBA1000213	16224	
HEMBA1000243	C-HEMBA1000243	16225	
HEMBA1000244	C-HEMBA1000244	16226	16227
HEMBA1000251	C-HEMBA1000251	16228	
HEMBA1000338	C-HEMBA1000338	16229	
HEMBA1000357	C-HEMBA1000357	16230	16231
HEMBA1000376	C-HEMBA1000376	16232	
HEMBA1000428	C-HEMBA1000428	16233	16234
HEMBA1000469	C-HEMBA1000469	16235	
HEMBA1000497	C-HEMBA1000497	16236	16237
HEMBA1000561	C-HEMBA1000561	16238	16239
HEMBA1000569	C-HEMBA1000569	16240	16241
HEMBA1000575	C-HEMBA1000575	16242	
HEMBA1000591	C-HEMBA1000591	16243	16244
HEMBA1000673	C-HEMBA1000673	16245	
HEMBA1000702	C-HEMBA1000702	16246	
HEMBA1000722	C-HEMBA1000722	16247	
HEMBA1000726	C-HEMBA1000726	16248	
HEMBA1000876	C-HEMBA1000876	16249	
HEMBA1000942	C-HEMBA1000942	16250	
HEMBA1000943	C-HEMBA1000943	16251	
HEMBA1000960	C-HEMBA1000960	16252	
HEMBA1000985	C-HEMBA1000985	16253	
HEMBA1001019	C-HEMBA1001019	16254	
HEMBA1001020	C-HEMBA1001020	16255	16256
HEMBA1001024	C-HEMBA1001024	16257	
HEMBA1001026	C-HEMBA1001026	16258	
HEMBA1001051	C-HEMBA1001051	16259	
HEMBA1001060	C-HEMBA1001060	16260	
HEMBA1001071	C-HEMBA1001071	16261	
HEMBA1001077	C-HEMBA1001077	16262	16263
HEMBA1001099	C-HEMBA1001099	16264	
HEMBA1001121	C-HEMBA1001121	16265	
HEMBA1001123	C-HEMBA1001123	16266	
HEMBA1001208	C-HEMBA1001208	16267	
HEMBA1001213	C-HEMBA1001213	16268	
HEMBA1001226	C-HEMBA1001226	16269	
HEMBA1001247	C-HEMBA1001247	16270	16271

(984)

特開 2002-191363

1965

1966

HEMBA1001299	C-HEMBA1001299	16272	
HEMBA1001319	C-HEMBA1001319	16273	
HEMBA1001323	C-HEMBA1001323	16274	
HEMBA1001327	C-HEMBA1001327	16275	
HEMBA1001361	C-HEMBA1001361	16276	
HEMBA1001375	C-HEMBA1001375	16277	
HEMBA1001377	C-HEMBA1001377	16278	
HEMBA1001383	C-HEMBA1001383	16279	
HEMBA1001391	C-HEMBA1001391	16280	
HEMBA1001411	C-HEMBA1001411	16281	
HEMBA1001432	C-HEMBA1001432	16282	
HEMBA1001433	C-HEMBA1001433	16283	
HEMBA1001435	C-HEMBA1001435	16284	
HEMBA1001442	C-HEMBA1001442	16285	
HEMBA1001463	C-HEMBA1001463	16286	
HEMBA1001515	C-HEMBA1001515	16287	16288
HEMBA1001522	C-HEMBA1001522	16289	16290
HEMBA1001557	C-HEMBA1001557	16291	16292
HEMBA1001566	C-HEMBA1001566	16293	
HEMBA1001589	C-HEMBA1001589	16294	
HEMBA1001608	C-HEMBA1001608	16295	16296
HEMBA1001636	C-HEMBA1001636	16297	16298
HEMBA1001647	C-HEMBA1001647	16299	16300
HEMBA1001651	C-HEMBA1001651	16301	16302
HEMBA1001658	C-HEMBA1001658	16303	
HEMBA1001675	C-HEMBA1001675	16304	16305
HEMBA1001712	C-HEMBA1001712	16306	
HEMBA1001734	C-HEMBA1001734	16307	
HEMBA1001745	C-HEMBA1001745	16308	
HEMBA1001750	C-HEMBA1001750	16309	
HEMBA1001784	C-HEMBA1001784	16310	16311
HEMBA1001791	C-HEMBA1001791	16312	16313
HEMBA1001803	C-HEMBA1001803	16314	
HEMBA1001820	C-HEMBA1001820	16315	
HEMBA1001835	C-HEMBA1001835	16316	
HEMBA1001888	C-HEMBA1001888	16317	
HEMBA1001912	C-HEMBA1001912	16318	
HEMBA1001915	C-HEMBA1001915	16319	
HEMBA1001918	C-HEMBA1001918	16320	
HEMBA1001940	C-HEMBA1001940	16321	
HEMBA1001942	C-HEMBA1001942	16322	
HEMBA1001964	C-HEMBA1001964	16323	
HEMBA1002022	C-HEMBA1002022	16324	
HEMBA1002039	C-HEMBA1002039	16325	16326
HEMBA1002100	C-HEMBA1002100	16327	
HEMBA1002113	C-HEMBA1002113	16328	16329
HEMBA1002119	C-HEMBA1002119	16330	
HEMBA1002139	C-HEMBA1002139	16331	
HEMBA1002160	C-HEMBA1002160	16332	
HEMBA1002162	C-HEMBA1002162	16333	

(985)

特開 2002-191363

1967

1968

HEMBA1002166	C-HEMBA1002166	16334	16335
HEMBA1002185	C-HEMBA1002185	16336	
HEMBA1002204	C-HEMBA1002204	16337	
HEMBA1002328	C-HEMBA1002328	16338	
HEMBA1002337	C-HEMBA1002337	16339	
HEMBA1002348	C-HEMBA1002348	16340	
HEMBA1002381	C-HEMBA1002381	16341	16342
HEMBA1002486	C-HEMBA1002486	16343	16344
HEMBA1002498	C-HEMBA1002498	16345	
HEMBA1002538	C-HEMBA1002538	16346	
HEMBA1002552	C-HEMBA1002552	16347	
HEMBA1002555	C-HEMBA1002555	16348	16349
HEMBA1002558	C-HEMBA1002558	16350	
HEMBA1002621	C-HEMBA1002621	16351	
HEMBA1002629	C-HEMBA1002629	16352	
HEMBA1002645	C-HEMBA1002645	16353	
HEMBA1002659	C-HEMBA1002659	16354	16355
HEMBA1002661	C-HEMBA1002661	16356	16357
HEMBA1002666	C-HEMBA1002666	16358	
HEMBA1002678	C-HEMBA1002678	16359	
HEMBA1002679	C-HEMBA1002679	16360	16361
HEMBA1002712	C-HEMBA1002712	16362	
HEMBA1002716	C-HEMBA1002716	16363	
HEMBA1002742	C-HEMBA1002742	16364	
HEMBA1002746	C-HEMBA1002746	16365	16366
HEMBA1002748	C-HEMBA1002748	16367	16368
HEMBA1002780	C-HEMBA1002780	16369	16370
HEMBA1002801	C-HEMBA1002801	16371	
HEMBA1002826	C-HEMBA1002826	16372	
HEMBA1002833	C-HEMBA1002833	16373	
HEMBA1002921	C-HEMBA1002921	16374	16375
HEMBA1002934	C-HEMBA1002934	16376	
HEMBA1002944	C-HEMBA1002944	16377	
HEMBA1002968	C-HEMBA1002968	16378	
HEMBA1003034	C-HEMBA1003034	16379	
HEMBA1003037	C-HEMBA1003037	16380	
HEMBA1003071	C-HEMBA1003071	16381	16382
HEMBA1003078	C-HEMBA1003078	16383	
HEMBA1003083	C-HEMBA1003083	16384	
HEMBA1003086	C-HEMBA1003086	16385	16386
HEMBA1003098	C-HEMBA1003098	16387	16388
HEMBA1003133	C-HEMBA1003133	16389	16390
HEMBA1003142	C-HEMBA1003142	16391	16392
HEMBA1003166	C-HEMBA1003166	16393	
HEMBA1003197	C-HEMBA1003197	16394	
HEMBA1003202	C-HEMBA1003202	16395	
HEMBA1003220	C-HEMBA1003220	16396	16397
HEMBA1003229	C-HEMBA1003229	16398	16399
HEMBA1003276	C-HEMBA1003276	16400	
HEMBA1003278	C-HEMBA1003278	16401	

(986)

特開 2002-191363

1969

1970

HEMBA1003328	C-HEMBA1003328	16402	
HEMBA1003373	C-HEMBA1003373	16403	
HEMBA1003597	C-HEMBA1003597	16404	16405
HEMBA1003598	C-HEMBA1003598	16406	16407
HEMBA1003656	C-HEMBA1003656	16408	
HEMBA1003680	C-HEMBA1003680	16409	16410
HEMBA1003733	C-HEMBA1003733	16411	
HEMBA1003742	C-HEMBA1003742	16412	16413
HEMBA1003760	C-HEMBA1003760	16414	16415
HEMBA1003803	C-HEMBA1003803	16416	16417
HEMBA1003854	C-HEMBA1003854	16418	16419
HEMBA1003926	C-HEMBA1003926	16420	
HEMBA1003939	C-HEMBA1003939	16421	
HEMBA1003987	C-HEMBA1003987	16422	
HEMBA1004012	C-HEMBA1004012	16423	
HEMBA1004015	C-HEMBA1004015	16424	
HEMBA1004193	C-HEMBA1004193	16425	
HEMBA1004225	C-HEMBA1004225	16426	
HEMBA1004241	C-HEMBA1004241	16427	
HEMBA1004267	C-HEMBA1004267	16428	
HEMBA1004295	C-HEMBA1004295	16429	16430
HEMBA1004354	C-HEMBA1004354	16431	16432
HEMBA1004356	C-HEMBA1004356	16433	16434
HEMBA1004396	C-HEMBA1004396	16435	
HEMBA1004405	C-HEMBA1004405	16436	
HEMBA1004433	C-HEMBA1004433	16437	
HEMBA1004538	C-HEMBA1004538	16438	16439
HEMBA1004542	C-HEMBA1004542	16440	
HEMBA1004573	C-HEMBA1004573	16441	16442
HEMBA1004577	C-HEMBA1004577	16443	16444
HEMBA1004604	C-HEMBA1004604	16445	16446
HEMBA1004617	C-HEMBA1004617	16447	
HEMBA1004631	C-HEMBA1004631	16448	
HEMBA1004705	C-HEMBA1004705	16449	
HEMBA1004733	C-HEMBA1004733	16450	
HEMBA1004748	C-HEMBA1004748	16451	
HEMBA1004778	C-HEMBA1004778	16452	
HEMBA1004803	C-HEMBA1004803	16453	
HEMBA1004807	C-HEMBA1004807	16454	
HEMBA1004820	C-HEMBA1004820	16455	
HEMBA1004865	C-HEMBA1004865	16456	
HEMBA1004880	C-HEMBA1004880	16457	
HEMBA1004900	C-HEMBA1004900	16458	
HEMBA1004909	C-HEMBA1004909	16459	
HEMBA1004960	C-HEMBA1004960	16460	
HEMBA1004978	C-HEMBA1004978	16461	
HEMBA1004980	C-HEMBA1004980	16462	16463
HEMBA1004983	C-HEMBA1004983	16464	
HEMBA1004995	C-HEMBA1004995	16465	16466
HEMBA1005019	C-HEMBA1005019	16467	16468

(987)

特開 2002-191363

1971

1972

HEMBA1005029	C-HEMBA1005029	16469	16470
HEMBA1005035	C-HEMBA1005035	16471	16472
HEMBA1005039	C-HEMBA1005039	16473	
HEMBA1005047	C-HEMBA1005047	16474	16475
HEMBA1005050	C-HEMBA1005050	16476	16477
HEMBA1005062	C-HEMBA1005062	16478	
HEMBA1005066	C-HEMBA1005066	16479	16480
HEMBA1005075	C-HEMBA1005075	16481	16482
HEMBA1005079	C-HEMBA1005079	16483	
HEMBA1005101	C-HEMBA1005101	16484	16485
HEMBA1005123	C-HEMBA1005123	16486	
HEMBA1005149	C-HEMBA1005149	16487	16488
HEMBA1005152	C-HEMBA1005152	16489	
HEMBA1005201	C-HEMBA1005201	16490	16491
HEMBA1005202	C-HEMBA1005202	16492	16493
HEMBA1005223	C-HEMBA1005223	16494	16495
HEMBA1005232	C-HEMBA1005232	16496	
HEMBA1005241	C-HEMBA1005241	16497	16498
HEMBA1005275	C-HEMBA1005275	16499	
HEMBA1005293	C-HEMBA1005293	16500	
HEMBA1005311	C-HEMBA1005311	16501	16502
HEMBA1005338	C-HEMBA1005338	16503	16504
HEMBA1005359	C-HEMBA1005359	16505	
HEMBA1005367	C-HEMBA1005367	16506	16507
HEMBA1005374	C-HEMBA1005374	16508	
HEMBA1005382	C-HEMBA1005382	16509	16510
HEMBA1005411	C-HEMBA1005411	16511	
HEMBA1005426	C-HEMBA1005426	16512	
HEMBA1005443	C-HEMBA1005443	16513	
HEMBA1005447	C-HEMBA1005447	16514	
HEMBA1005497	C-HEMBA1005497	16515	
HEMBA1005500	C-HEMBA1005500	16516	
HEMBA1005506	C-HEMBA1005506	16517	
HEMBA1005508	C-HEMBA1005508	16518	
HEMBA1005526	C-HEMBA1005526	16519	
HEMBA1005530	C-HEMBA1005530	16520	16521
HEMBA1005548	C-HEMBA1005548	16522	16523
HEMBA1005552	C-HEMBA1005552	16524	16525
HEMBA1005568	C-HEMBA1005568	16526	
HEMBA1005588	C-HEMBA1005588	16527	16528
HEMBA1005593	C-HEMBA1005593	16529	16530
HEMBA1005606	C-HEMBA1005606	16531	
HEMBA1005616	C-HEMBA1005616	16532	16533
HEMBA1005627	C-HEMBA1005627	16534	
HEMBA1005670	C-HEMBA1005670	16535	
HEMBA1005679	C-HEMBA1005679	16536	
HEMBA1005699	C-HEMBA1005699	16537	16538
HEMBA1005705	C-HEMBA1005705	16539	
HEMBA1005732	C-HEMBA1005732	16540	16541
HEMBA1005815	C-HEMBA1005815	16542	16543

(988)

特開 2002-191363

1973

1974

HEMBA1005852	C-HEMBA1005852	16544	16545
HEMBA1005894	C-HEMBA1005894	16546	
HEMBA1005921	C-HEMBA1005921	16547	
HEMBA1006035	C-HEMBA1006035	16548	16549
HEMBA1006036	C-HEMBA1006036	16550	16551
HEMBA1006090	C-HEMBA1006090	16552	
HEMBA1006138	C-HEMBA1006138	16553	16554
HEMBA1006173	C-HEMBA1006173	16555	16556
HEMBA1006252	C-HEMBA1006252	16557	16558
HEMBA1006268	C-HEMBA1006268	16559	16560
HEMBA1006347	C-HEMBA1006347	16561	16562
HEMBA1006359	C-HEMBA1006359	16563	
HEMBA1006380	C-HEMBA1006380	16564	
HEMBA1006416	C-HEMBA1006416	16565	16566
HEMBA1006421	C-HEMBA1006421	16567	
HEMBA1006424	C-HEMBA1006424	16568	
HEMBA1006426	C-HEMBA1006426	16569	
HEMBA1006446	C-HEMBA1006446	16570	
HEMBA1006485	C-HEMBA1006485	16571	16572
HEMBA1006486	C-HEMBA1006486	16573	
HEMBA1006494	C-HEMBA1006494	16574	
HEMBA1006546	C-HEMBA1006546	16575	
HEMBA1006562	C-HEMBA1006562	16576	
HEMBA1006595	C-HEMBA1006595	16577	
HEMBA1006597	C-HEMBA1006597	16578	
HEMBA1006631	C-HEMBA1006631	16579	
HEMBA1006639	C-HEMBA1006639	16580	
HEMBA1006652	C-HEMBA1006652	16581	16582
HEMBA1006659	C-HEMBA1006659	16583	16584
HEMBA1006665	C-HEMBA1006665	16585	
HEMBA1006676	C-HEMBA1006676	16586	16587
HEMBA1006695	C-HEMBA1006695	16588	
HEMBA1006709	C-HEMBA1006709	16589	16590
HEMBA1006758	C-HEMBA1006758	16591	16592
HEMBA1006780	C-HEMBA1006780	16593	
HEMBA1006807	C-HEMBA1006807	16594	16595
HEMBA1006824	C-HEMBA1006824	16596	
HEMBA1006865	C-HEMBA1006865	16597	
HEMBA1006921	C-HEMBA1006921	16598	16599
HEMBA1006949	C-HEMBA1006949	16600	
HEMBA1006976	C-HEMBA1006976	16601	16602
HEMBA1007051	C-HEMBA1007051	16603	
HEMBA1007052	C-HEMBA1007052	16604	16605
HEMBA1007066	C-HEMBA1007066	16606	
HEMBA1007073	C-HEMBA1007073	16607	
HEMBA1007078	C-HEMBA1007078	16608	
HEMBA1007085	C-HEMBA1007085	16609	16610
HEMBA1007113	C-HEMBA1007113	16611	
HEMBA1007121	C-HEMBA1007121	16612	
HEMBA1007129	C-HEMBA1007129	16613	

(989)

特開 2002-191363

1975

1976

HEMBA1007147	C-HEMBA1007147	16614	
HEMBA1007151	C-HEMBA1007151	16615	16616
HEMBA1007178	C-HEMBA1007178	16617	
HEMBA1007203	C-HEMBA1007203	16618	16619
HEMBA1007224	C-HEMBA1007224	16620	16621
HEMBA1007243	C-HEMBA1007243	16622	16623
HEMBA1007251	C-HEMBA1007251	16624	16625
HEMBA1007288	C-HEMBA1007288	16626	
HEMBA1007322	C-HEMBA1007322	16627	
HEMBA1007341	C-HEMBA1007341	16628	
HEMBA1000050	C-HEMBA1000050	16629	16630
HEMBA1000054	C-HEMBA1000054	16631	16632
HEMBA1000059	C-HEMBA1000059	16633	16634
HEMBA1000089	C-HEMBA1000089	16635	
HEMBA1000113	C-HEMBA1000113	16636	
HEMBA1000144	C-HEMBA1000144	16637	16638
HEMBA1000173	C-HEMBA1000173	16639	
HEMBA1000175	C-HEMBA1000175	16640	16641
HEMBA1000272	C-HEMBA1000272	16642	
HEMBA1000317	C-HEMBA1000317	16643	16644
HEMBA1000318	C-HEMBA1000318	16645	
HEMBA1000336	C-HEMBA1000336	16646	
HEMBA1000341	C-HEMBA1000341	16647	
HEMBA1000343	C-HEMBA1000343	16648	
HEMBA1000354	C-HEMBA1000354	16649	
HEMBA1000374	C-HEMBA1000374	16650	
HEMBA1000434	C-HEMBA1000434	16651	
HEMBA1000441	C-HEMBA1000441	16652	
HEMBA1000491	C-HEMBA1000491	16653	
HEMBA1000493	C-HEMBA1000493	16654	
HEMBA1000510	C-HEMBA1000510	16655	
HEMBA1000652	C-HEMBA1000652	16656	
HEMBA1000672	C-HEMBA1000672	16657	
HEMBA1000684	C-HEMBA1000684	16658	
HEMBA1000709	C-HEMBA1000709	16659	
HEMBA1000726	C-HEMBA1000726	16660	
HEMBA1000770	C-HEMBA1000770	16661	
HEMBA1000827	C-HEMBA1000827	16662	16663
HEMBA1000831	C-HEMBA1000831	16664	16665
HEMBA1000883	C-HEMBA1000883	16666	
HEMBA1000888	C-HEMBA1000888	16667	
HEMBA1000893	C-HEMBA1000893	16668	
HEMBA1000913	C-HEMBA1000913	16669	
HEMBA1000996	C-HEMBA1000996	16670	
HEMBA1001004	C-HEMBA1001004	16671	
HEMBA1001047	C-HEMBA1001047	16672	
HEMBA1001060	C-HEMBA1001060	16673	16674
HEMBA1001114	C-HEMBA1001114	16675	
HEMBA1001119	C-HEMBA1001119	16676	
HEMBA1001133	C-HEMBA1001133	16677	

(990)

特開 2002-191363

1977

1978

HEM BB1001142	C-HEM BB1001142	16678	
HEM BB1001177	C-HEM BB1001177	16679	16680
HEM BB1001208	C-HEM BB1001208	16681	
HEM BB1001209	C-HEM BB1001209	16682	
HEM BB1001249	C-HEM BB1001249	16683	
HEM BB1001253	C-HEM BB1001253	16684	16685
HEM BB1001254	C-HEM BB1001254	16686	
HEM BB1001271	C-HEM BB1001271	16687	16688
HEM BB1001304	C-HEM BB1001304	16689	16690
HEM BB1001317	C-HEM BB1001317	16691	
HEM BB1001348	C-HEM BB1001348	16692	16693
HEM BB1001394	C-HEM BB1001394	16694	
HEM BB1001410	C-HEM BB1001410	16695	
HEM BB1001424	C-HEM BB1001424	16696	
HEM BB1001426	C-HEM BB1001426	16697	
HEM BB1001429	C-HEM BB1001429	16698	16699
HEM BB1001436	C-HEM BB1001436	16700	
HEM BB1001443	C-HEM BB1001443	16701	16702
HEM BB1001449	C-HEM BB1001449	16703	16704
HEM BB1001458	C-HEM BB1001458	16705	
HEM BB1001521	C-HEM BB1001521	16706	
HEM BB1001531	C-HEM BB1001531	16707	16708
HEM BB1001535	C-HEM BB1001535	16709	
HEM BB1001536	C-HEM BB1001536	16710	
HEM BB1001564	C-HEM BB1001564	16711	16712
HEM BB1001565	C-HEM BB1001565	16713	16714
HEM BB1001585	C-HEM BB1001585	16715	
HEM BB1001588	C-HEM BB1001588	16716	
HEM BB1001603	C-HEM BB1001603	16717	
HEM BB1001618	C-HEM BB1001618	16718	
HEM BB1001635	C-HEM BB1001635	16719	
HEM BB1001653	C-HEM BB1001653	16720	
HEM BB1001668	C-HEM BB1001668	16721	
HEM BB1001673	C-HEM BB1001673	16722	16723
HEM BB1001685	C-HEM BB1001685	16724	
HEM BB1001695	C-HEM BB1001695	16725	16726
HEM BB1001707	C-HEM BB1001707	16727	16728
HEM BB1001735	C-HEM BB1001735	16729	16730
HEM BB1001736	C-HEM BB1001736	16731	16732
HEM BB1001747	C-HEM BB1001747	16733	
HEM BB1001749	C-HEM BB1001749	16734	
HEM BB1001753	C-HEM BB1001753	16735	
HEM BB1001756	C-HEM BB1001756	16736	
HEM BB1001760	C-HEM BB1001760	16737	
HEM BB1001785	C-HEM BB1001785	16738	
HEM BB1001797	C-HEM BB1001797	16739	
HEM BB1001802	C-HEM BB1001802	16740	16741
HEM BB1001816	C-HEM BB1001816	16742	16743
HEM BB1001831	C-HEM BB1001831	16744	16745
HEM BB1001839	C-HEM BB1001839	16746	16747

(991)

特開 2002-191363

1979

1980

HEMBB1001850	C-HEMBB1001850	16748	
HEMBB1001863	C-HEMBB1001863	16749	
HEMBB1001868	C-HEMBB1001868	16750	
HEMBB1001874	C-HEMBB1001874	16751	16752
HEMBB1001880	C-HEMBB1001880	16753	
HEMBB1001899	C-HEMBB1001899	16754	
HEMBB1001906	C-HEMBB1001906	16755	
HEMBB1001910	C-HEMBB1001910	16756	
HEMBB1001911	C-HEMBB1001911	16757	
HEMBB1001921	C-HEMBB1001921	16758	
HEMBB1001922	C-HEMBB1001922	16759	
HEMBB1001930	C-HEMBB1001930	16760	
HEMBB1001944	C-HEMBB1001944	16761	
HEMBB1001945	C-HEMBB1001945	16762	16763
HEMBB1001947	C-HEMBB1001947	16764	16765
HEMBB1001950	C-HEMBB1001950	16766	16767
HEMBB1001952	C-HEMBB1001952	16768	
HEMBB1001957	C-HEMBB1001957	16769	
HEMBB1001962	C-HEMBB1001962	16770	16771
HEMBB1001983	C-HEMBB1001983	16772	
HEMBB1001990	C-HEMBB1001990	16773	
HEMBB1001996	C-HEMBB1001996	16774	
HEMBB1002002	C-HEMBB1002002	16775	16776
HEMBB1002005	C-HEMBB1002005	16777	
HEMBB1002042	C-HEMBB1002042	16778	
HEMBB1002043	C-HEMBB1002043	16779	
HEMBB1002045	C-HEMBB1002045	16780	16781
HEMBB1002049	C-HEMBB1002049	16782	
HEMBB1002050	C-HEMBB1002050	16783	16784
HEMBB1002068	C-HEMBB1002068	16785	16786
HEMBB1002092	C-HEMBB1002092	16787	16788
HEMBB1002139	C-HEMBB1002139	16789	
HEMBB1002142	C-HEMBB1002142	16790	16791
HEMBB1002190	C-HEMBB1002190	16792	16793
HEMBB1002193	C-HEMBB1002193	16794	16795
HEMBB1002217	C-HEMBB1002217	16796	16797
HEMBB1002218	C-HEMBB1002218	16798	
HEMBB1002232	C-HEMBB1002232	16799	
HEMBB1002247	C-HEMBB1002247	16800	16801
HEMBB1002249	C-HEMBB1002249	16802	
HEMBB1002266	C-HEMBB1002266	16803	16804
HEMBB1002327	C-HEMBB1002327	16805	
HEMBB1002329	C-HEMBB1002329	16806	16807
HEMBB1002342	C-HEMBB1002342	16808	16809
HEMBB1002358	C-HEMBB1002358	16810	
HEMBB1002371	C-HEMBB1002371	16811	
HEMBB1002387	C-HEMBB1002387	16812	16813
HEMBB1002409	C-HEMBB1002409	16814	16815
HEMBB1002425	C-HEMBB1002425	16816	
HEMBB1002442	C-HEMBB1002442	16817	16818

(992)

特開 2002-191363

1981

1982

HEMBB1002453	C-HEMBB1002453	16819	
HEMBB1002458	C-HEMBB1002458	16820	
HEMBB1002477	C-HEMBB1002477	16821	16822
HEMBB1002489	C-HEMBB1002489	16823	
HEMBB1002510	C-HEMBB1002510	16824	16825
HEMBB1002520	C-HEMBB1002520	16826	
HEMBB1002522	C-HEMBB1002522	16827	
HEMBB1002545	C-HEMBB1002545	16828	
HEMBB1002579	C-HEMBB1002579	16829	
HEMBB1002582	C-HEMBB1002582	16830	
HEMBB1002596	C-HEMBB1002596	16831	
HEMBB1002603	C-HEMBB1002603	16832	
HEMBB1002610	C-HEMBB1002610	16833	
HEMBB1002613	C-HEMBB1002613	16834	
HEMBB1002617	C-HEMBB1002617	16835	
HEMBB1002623	C-HEMBB1002623	16836	
HEMBB1002635	C-HEMBB1002635	16837	
HEMBB1002677	C-HEMBB1002677	16838	
HEMBB1002683	C-HEMBB1002683	16839	
HEMBB1002699	C-HEMBB1002699	16840	
HEMBB1002702	C-HEMBB1002702	16841	
MAMMA1000009	C-MAMMA1000009	16842	
MAMMA1000043	C-MAMMA1000043	16843	
MAMMA1000045	C-MAMMA1000045	16844	16845
MAMMA1000057	C-MAMMA1000057	16846	16847
MAMMA1000085	C-MAMMA1000085	16848	16849
MAMMA1000092	C-MAMMA1000092	16850	
MAMMA1000103	C-MAMMA1000103	16851	
MAMMA1000117	C-MAMMA1000117	16852	16853
MAMMA1000129	C-MAMMA1000129	16854	16855
MAMMA1000133	C-MAMMA1000133	16856	16857
MAMMA1000155	C-MAMMA1000155	16858	
MAMMA1000175	C-MAMMA1000175	16859	
MAMMA1000198	C-MAMMA1000198	16860	
MAMMA1000241	C-MAMMA1000241	16861	
MAMMA1000251	C-MAMMA1000251	16862	16863
MAMMA1000254	C-MAMMA1000254	16864	16865
MAMMA1000287	C-MAMMA1000287	16866	
MAMMA1000307	C-MAMMA1000307	16867	
MAMMA1000331	C-MAMMA1000331	16868	
MAMMA1000339	C-MAMMA1000339	16869	16870
MAMMA1000340	C-MAMMA1000340	16871	
MAMMA1000348	C-MAMMA1000348	16872	
MAMMA1000356	C-MAMMA1000356	16873	
MAMMA1000360	C-MAMMA1000360	16874	
MAMMA1000402	C-MAMMA1000402	16875	16876
MAMMA1000414	C-MAMMA1000414	16877	
MAMMA1000431	C-MAMMA1000431	16878	16879
MAMMA1000444	C-MAMMA1000444	16880	
MAMMA1000458	C-MAMMA1000458	16881	

(993)

特開 2002-191363

1983

1984

MAMMA1000500	C-MAMMA1000500	16882	
MAMMA1000522	C-MAMMA1000522	16883	16884
MAMMA1000576	C-MAMMA1000576	16885	
MAMMA1000583	C-MAMMA1000583	16886	
MAMMA1000594	C-MAMMA1000594	16887	
MAMMA1000605	C-MAMMA1000605	16888	
MAMMA1000616	C-MAMMA1000616	16889	
MAMMA1000643	C-MAMMA1000643	16890	
MAMMA1000684	C-MAMMA1000684	16891	16892
MAMMA1000696	C-MAMMA1000696	16893	
MAMMA1000707	C-MAMMA1000707	16894	
MAMMA1000714	C-MAMMA1000714	16895	16896
MAMMA1000720	C-MAMMA1000720	16897	
MAMMA1000744	C-MAMMA1000744	16898	
MAMMA1000761	C-MAMMA1000761	16899	
MAMMA1000776	C-MAMMA1000776	16900	16901
MAMMA1000798	C-MAMMA1000798	16902	16903
MAMMA1000839	C-MAMMA1000839	16904	
MAMMA1000851	C-MAMMA1000851	16905	
MAMMA1000863	C-MAMMA1000863	16906	
MAMMA1000867	C-MAMMA1000867	16907	
MAMMA1000876	C-MAMMA1000876	16908	16909
MAMMA1000880	C-MAMMA1000880	16910	
MAMMA1000883	C-MAMMA1000883	16911	
MAMMA1000921	C-MAMMA1000921	16912	
MAMMA1000931	C-MAMMA1000931	16913	
MAMMA1000941	C-MAMMA1000941	16914	
MAMMA1000957	C-MAMMA1000957	16915	
MAMMA1000962	C-MAMMA1000962	16916	
MAMMA1000975	C-MAMMA1000975	16917	
MAMMA1000987	C-MAMMA1000987	16918	
MAMMA1001003	C-MAMMA1001003	16919	
MAMMA1001030	C-MAMMA1001030	16920	16921
MAMMA1001038	C-MAMMA1001038	16922	
MAMMA1001082	C-MAMMA1001082	16923	
MAMMA1001162	C-MAMMA1001162	16924	16925
MAMMA1001186	C-MAMMA1001186	16926	
MAMMA1001191	C-MAMMA1001191	16927	16928
MAMMA1001206	C-MAMMA1001206	16929	
MAMMA1001220	C-MAMMA1001220	16930	
MAMMA1001243	C-MAMMA1001243	16931	
MAMMA1001249	C-MAMMA1001249	16932	
MAMMA1001256	C-MAMMA1001256	16933	
MAMMA1001268	C-MAMMA1001268	16934	16935
MAMMA1001271	C-MAMMA1001271	16936	16937
MAMMA1001274	C-MAMMA1001274	16938	16939
MAMMA1001292	C-MAMMA1001292	16940	16941
MAMMA1001305	C-MAMMA1001305	16942	16943
MAMMA1001324	C-MAMMA1001324	16944	
MAMMA1001341	C-MAMMA1001341	16945	

(994)

特開 2002-191363

1985

1986

MAMMA1001388	C-MAMMA1001388	16946	16947
MAMMA1001397	C-MAMMA1001397	16948	
MAMMA1001408	C-MAMMA1001408	16949	
MAMMA1001420	C-MAMMA1001420	16950	
MAMMA1001442	C-MAMMA1001442	16951	
MAMMA1001452	C-MAMMA1001452	16952	
MAMMA1001465	C-MAMMA1001465	16953	16954
MAMMA1001487	C-MAMMA1001487	16955	16956
MAMMA1001501	C-MAMMA1001501	16957	
MAMMA1001547	C-MAMMA1001547	16958	
MAMMA1001551	C-MAMMA1001551	16959	16960
MAMMA1001575	C-MAMMA1001575	16961	
MAMMA1001590	C-MAMMA1001590	16962	16963
MAMMA1001600	C-MAMMA1001600	16964	16965
MAMMA1001606	C-MAMMA1001606	16966	
MAMMA1001627	C-MAMMA1001627	16967	16968
MAMMA1001663	C-MAMMA1001663	16969	
MAMMA1001670	C-MAMMA1001670	16970	16971
MAMMA1001671	C-MAMMA1001671	16972	
MAMMA1001679	C-MAMMA1001679	16973	
MAMMA1001711	C-MAMMA1001711	16974	
MAMMA1001735	C-MAMMA1001735	16975	16976
MAMMA1001744	C-MAMMA1001744	16977	
MAMMA1001745	C-MAMMA1001745	16978	
MAMMA1001751	C-MAMMA1001751	16979	16980
MAMMA1001783	C-MAMMA1001783	16981	
MAMMA1001788	C-MAMMA1001788	16982	
MAMMA1001806	C-MAMMA1001806	16983	
MAMMA1001812	C-MAMMA1001812	16984	
MAMMA1001815	C-MAMMA1001815	16985	
MAMMA1001817	C-MAMMA1001817	16986	16987
MAMMA1001818	C-MAMMA1001818	16988	
MAMMA1001820	C-MAMMA1001820	16989	16990
MAMMA1001824	C-MAMMA1001824	16991	
MAMMA1001851	C-MAMMA1001851	16992	
MAMMA1001854	C-MAMMA1001854	16993	
MAMMA1001864	C-MAMMA1001864	16994	
MAMMA1001878	C-MAMMA1001878	16995	
MAMMA1001890	C-MAMMA1001890	16996	
MAMMA1001907	C-MAMMA1001907	16997	
MAMMA1001908	C-MAMMA1001908	16998	16999
MAMMA1001931	C-MAMMA1001931	17000	
MAMMA1001969	C-MAMMA1001969	17001	
MAMMA1002011	C-MAMMA1002011	17002	17003
MAMMA1002032	C-MAMMA1002032	17004	
MAMMA1002041	C-MAMMA1002041	17005	
MAMMA1002047	C-MAMMA1002047	17006	
MAMMA1002056	C-MAMMA1002056	17007	
MAMMA1002058	C-MAMMA1002058	17008	
MAMMA1002078	C-MAMMA1002078	17009	

(995)

特開 2002-191363

1987

1988

MAMMA1002082	C-MAMMA1002082	17010	
MAMMA1002084	C-MAMMA1002084	17011	
MAMMA1002093	C-MAMMA1002093	17012	
MAMMA1002094	C-MAMMA1002094	17013	
MAMMA1002118	C-MAMMA1002118	17014	
MAMMA1002125	C-MAMMA1002125	17015	
MAMMA1002132	C-MAMMA1002132	17016	
MAMMA1002140	C-MAMMA1002140	17017	
MAMMA1002143	C-MAMMA1002143	17018	17019
MAMMA1002145	C-MAMMA1002145	17020	
MAMMA1002198	C-MAMMA1002198	17021	
MAMMA1002230	C-MAMMA1002230	17022	
MAMMA1002250	C-MAMMA1002250	17023	
MAMMA1002282	C-MAMMA1002282	17024	17025
MAMMA1002293	C-MAMMA1002293	17026	
MAMMA1002298	C-MAMMA1002298	17027	17028
MAMMA1002299	C-MAMMA1002299	17029	
MAMMA1002308	C-MAMMA1002308	17030	
MAMMA1002310	C-MAMMA1002310	17031	17032
MAMMA1002311	C-MAMMA1002311	17033	
MAMMA1002322	C-MAMMA1002322	17034	
MAMMA1002339	C-MAMMA1002339	17035	
MAMMA1002352	C-MAMMA1002352	17036	
MAMMA1002359	C-MAMMA1002359	17037	17038
MAMMA1002360	C-MAMMA1002360	17039	17040
MAMMA1002392	C-MAMMA1002392	17041	17042
MAMMA1002411	C-MAMMA1002411	17043	
MAMMA1002413	C-MAMMA1002413	17044	
MAMMA1002417	C-MAMMA1002417	17045	
MAMMA1002428	C-MAMMA1002428	17046	
MAMMA1002434	C-MAMMA1002434	17047	
MAMMA1002446	C-MAMMA1002446	17048	
MAMMA1002454	C-MAMMA1002454	17049	
MAMMA1002461	C-MAMMA1002461	17050	17051
MAMMA1002475	C-MAMMA1002475	17052	17053
MAMMA1002556	C-MAMMA1002556	17054	
MAMMA1002566	C-MAMMA1002566	17055	
MAMMA1002612	C-MAMMA1002612	17056	
MAMMA1002622	C-MAMMA1002622	17057	17058
MAMMA1002637	C-MAMMA1002637	17059	17060
MAMMA1002650	C-MAMMA1002650	17061	17062
MAMMA1002699	C-MAMMA1002699	17063	17064
MAMMA1002727	C-MAMMA1002727	17065	
MAMMA1002748	C-MAMMA1002748	17066	
MAMMA1002758	C-MAMMA1002758	17067	
MAMMA1002780	C-MAMMA1002780	17068	
MAMMA1002820	C-MAMMA1002820	17069	
MAMMA1002833	C-MAMMA1002833	17070	
MAMMA1002843	C-MAMMA1002843	17071	17072
MAMMA1002895	C-MAMMA1002895	17073	17074

(996)

特開 2002-191363

1989

1990

MAMMA1002937	C-MAMMA1002937	17075	17076
MAMMA1003004	C-MAMMA1003004	17077	
MAMMA1003047	C-MAMMA1003047	17078	17079
NT2RM1000001	C-NT2RM1000001	17080	17081
NT2RM1000018	C-NT2RM1000018	17082	17083
NT2RM1000037	C-NT2RM1000037	17084	17085
NT2RM1000086	C-NT2RM1000086	17086	17087
NT2RM1000421	C-NT2RM1000421	17088	17089
NT2RM1000499	C-NT2RM1000499	17090	17091
NT2RM1001059	C-NT2RM1001059	17092	17093
NT2RM1001092	C-NT2RM1001092	17094	17095
NT2RM2001592	C-NT2RM2001592	17096	17097
NT2RM2001635	C-NT2RM2001635	17098	17099
NT2RM2001637	C-NT2RM2001637	17100	17101
NT2RM2001641	C-NT2RM2001641	17102	
NT2RM2001670	C-NT2RM2001670	17103	17104
NT2RM2001699	C-NT2RM2001699	17105	17106
NT2RM2001706	C-NT2RM2001706	17107	
NT2RM2001718	C-NT2RM2001718	17108	17109
NT2RM2001727	C-NT2RM2001727	17110	17111
NT2RM2001805	C-NT2RM2001805	17112	17113
NT2RM4000086	C-NT2RM4000086	17114	17115
NT2RM4000215	C-NT2RM4000215	17116	17117
NT2RM4000414	C-NT2RM4000414	17118	
NT2RM4000590	C-NT2RM4000590	17119	17120
NT2RM4000634	C-NT2RM4000634	17121	17122
NT2RM4000657	C-NT2RM4000657	17123	17124
NT2RM4000783	C-NT2RM4000783	17125	
NT2RM4000857	C-NT2RM4000857	17126	17127
NT2RM4000971	C-NT2RM4000971	17128	17129
NT2RM4000996	C-NT2RM4000996	17130	17131
NT2RM4001092	C-NT2RM4001092	17132	17133
NT2RM4001178	C-NT2RM4001178	17134	17135
NT2RM4001569	C-NT2RM4001569	17136	17137
NT2RM4001819	C-NT2RM4001819	17138	
NT2RM4001905	C-NT2RM4001905	17139	17140
NT2RM4001938	C-NT2RM4001938	17141	17142
NT2RM4002062	C-NT2RM4002062	17143	17144
NT2RM4002073	C-NT2RM4002073	17145	17146
NT2RM4002093	C-NT2RM4002093	17147	17148
NT2RM4002109	C-NT2RM4002109	17149	17150
NT2RM4002146	C-NT2RM4002146	17151	17152
NT2RM4002194	C-NT2RM4002194	17153	17154
NT2RM4002390	C-NT2RM4002390	17155	17156
NT2RM4002398	C-NT2RM4002398	17157	17158
NT2RM4002420	C-NT2RM4002420	17159	
NT2RM4002534	C-NT2RM4002534	17160	17161
NT2RM4002565	C-NT2RM4002565	17162	17163
NT2RM4002571	C-NT2RM4002571	17164	17165
NT2RP1000358	C-NT2RP1000358	17166	17167

(997)

特開 2002-191363

1991

1992

NT2RP1000522	C-NT2RP1000522	17168	17169
NT2RP1000609	C-NT2RP1000609	17170	17171
NT2RP1000677	C-NT2RP1000677	17172	17173
NT2RP1000701	C-NT2RP1000701	17174	17175
NT2RP1000834	C-NT2RP1000834	17176	17177
NT2RP1000860	C-NT2RP1000860	17178	17179
NT2RP1000916	C-NT2RP1000916	17180	17181
NT2RP1000944	C-NT2RP1000944	17182	
NT2RP1001079	C-NT2RP1001079	17183	17184
NT2RP1001080	C-NT2RP1001080	17185	17186
NT2RP1001113	C-NT2RP1001113	17187	17188
NT2RP1001173	C-NT2RP1001173	17189	
NT2RP1001177	C-NT2RP1001177	17190	17191
NT2RP1001185	C-NT2RP1001185	17192	
NT2RP1001247	C-NT2RP1001247	17193	17194
NT2RP1001311	C-NT2RP1001311	17195	17196
NT2RP1001313	C-NT2RP1001313	17197	17198
NT2RP2000001	C-NT2RP2000001	17199	17200
NT2RP2000027	C-NT2RP2000027	17201	
NT2RP2000183	C-NT2RP2000183	17202	17203
NT2RP2000198	C-NT2RP2000198	17204	17205
NT2RP2000523	C-NT2RP2000523	17206	17207
NT2RP2000551	C-NT2RP2000551	17208	
NT2RP2000644	C-NT2RP2000644	17209	
NT2RP2000660	C-NT2RP2000660	17210	17211
NT2RP2000678	C-NT2RP2000678	17212	
NT2RP2000715	C-NT2RP2000715	17213	
NT2RP2000842	C-NT2RP2000842	17214	17215
NT2RP2000970	C-NT2RP2000970	17216	17217
NT2RP2001347	C-NT2RP2001347	17218	17219
NT2RP2001460	C-NT2RP2001460	17220	17221
NT2RP2001613	C-NT2RP2001613	17222	17223
NT2RP2001634	C-NT2RP2001634	17224	17225
NT2RP2001660	C-NT2RP2001660	17226	17227
NT2RP2001677	C-NT2RP2001677	17228	17229
NT2RP2001678	C-NT2RP2001678	17230	17231
NT2RP2001720	C-NT2RP2001720	17232	17233
NT2RP2001740	C-NT2RP2001740	17234	17235
NT2RP2001756	C-NT2RP2001756	17236	17237
NT2RP2001839	C-NT2RP2001839	17238	17239
NT2RP2001861	C-NT2RP2001861	17240	17241
NT2RP2001869	C-NT2RP2001869	17242	17243
NT2RP2001876	C-NT2RP2001876	17244	17245
NT2RP2001898	C-NT2RP2001898	17246	17247
NT2RP2001936	C-NT2RP2001936	17248	
NT2RP2001943	C-NT2RP2001943	17249	17250
NT2RP2001946	C-NT2RP2001946	17251	17252
NT2RP2002032	C-NT2RP2002032	17253	17254
NT2RP2002033	C-NT2RP2002033	17255	
NT2RP2002041	C-NT2RP2002041	17256	17257

(998)

特開 2002-191363

1993

1994

NT2RP2002047	C-NT2RP2002047	17258	
NT2RP2002066	C-NT2RP2002066	17259	17260
NT2RP2002124	C-NT2RP2002124	17261	17262
NT2RP2002172	C-NT2RP2002172	17263	
NT2RP2002219	C-NT2RP2002219	17264	
NT2RP2002256	C-NT2RP2002256	17265	17266
NT2RP2002316	C-NT2RP2002316	17267	
NT2RP2002373	C-NT2RP2002373	17268	17269
NT2RP2002439	C-NT2RP2002439	17270	17271
NT2RP2002475	C-NT2RP2002475	17272	17273
NT2RP2002546	C-NT2RP2002546	17274	
NT2RP2002591	C-NT2RP2002591	17275	17276
NT2RP2002606	C-NT2RP2002606	17277	17278
NT2RP2002643	C-NT2RP2002643	17279	17280
NT2RP2002727	C-NT2RP2002727	17281	17282
NT2RP2002736	C-NT2RP2002736	17283	17284
NT2RP2002740	C-NT2RP2002740	17285	17286
NT2RP2002741	C-NT2RP2002741	17287	17288
NT2RP2002752	C-NT2RP2002752	17289	
NT2RP2002753	C-NT2RP2002753	17290	17291
NT2RP2002857	C-NT2RP2002857	17292	17293
NT2RP2003000	C-NT2RP2003000	17294	17295
NT2RP2003073	C-NT2RP2003073	17296	
NT2RP2003164	C-NT2RP2003164	17297	17298
NT2RP2003206	C-NT2RP2003206	17299	
NT2RP2003228	C-NT2RP2003228	17300	17301
NT2RP2003230	C-NT2RP2003230	17302	17303
NT2RP2003237	C-NT2RP2003237	17304	
NT2RP2003272	C-NT2RP2003272	17305	17306
NT2RP2003280	C-NT2RP2003280	17307	17308
NT2RP2003293	C-NT2RP2003293	17309	
NT2RP2003394	C-NT2RP2003394	17310	17311
NT2RP2003401	C-NT2RP2003401	17312	17313
NT2RP2003456	C-NT2RP2003456	17314	
NT2RP2003517	C-NT2RP2003517	17315	
NT2RP2003522	C-NT2RP2003522	17316	17317
NT2RP2003559	C-NT2RP2003559	17318	17319
NT2RP2003564	C-NT2RP2003564	17320	17321
NT2RP2003581	C-NT2RP2003581	17322	17323
NT2RP2003643	C-NT2RP2003643	17324	17325
NT2RP2003702	C-NT2RP2003702	17326	17327
NT2RP2003704	C-NT2RP2003704	17328	17329
NT2RP2003727	C-NT2RP2003727	17330	17331
NT2RP2003751	C-NT2RP2003751	17332	17333
NT2RP2003781	C-NT2RP2003781	17334	17335
NT2RP2003825	C-NT2RP2003825	17336	17337
NT2RP2003871	C-NT2RP2003871	17338	17339
NT2RP2003885	C-NT2RP2003885	17340	
NT2RP2003912	C-NT2RP2003912	17341	17342
NT2RP2003976	C-NT2RP2003976	17343	

(999)

特開 2002-191363

1995

1996

NT2RP2003988	C-NT2RP2003988	17344	
NT2RP2004013	C-NT2RP2004013	17345	17346
NT2RP2004098	C-NT2RP2004098	17347	17348
NT2RP2004142	C-NT2RP2004142	17349	17350
NT2RP2004170	C-NT2RP2004170	17351	17352
NT2RP2004194	C-NT2RP2004194	17353	17354
NT2RP2004207	C-NT2RP2004207	17355	
NT2RP2004226	C-NT2RP2004226	17356	17357
NT2RP2004232	C-NT2RP2004232	17358	17359
NT2RP2004242	C-NT2RP2004242	17360	17361
NT2RP2004270	C-NT2RP2004270	17362	17363
NT2RP2004321	C-NT2RP2004321	17364	
NT2RP2004339	C-NT2RP2004339	17365	
NT2RP2004347	C-NT2RP2004347	17366	17367
NT2RP2004396	C-NT2RP2004396	17368	17369
NT2RP2004399	C-NT2RP2004399	17370	
NT2RP2004400	C-NT2RP2004400	17371	
NT2RP2004412	C-NT2RP2004412	17372	17373
NT2RP2004425	C-NT2RP2004425	17374	17375
NT2RP2004490	C-NT2RP2004490	17376	
NT2RP2004523	C-NT2RP2004523	17377	
NT2RP2004538	C-NT2RP2004538	17378	17379
NT2RP2004580	C-NT2RP2004580	17380	
NT2RP2004587	C-NT2RP2004587	17381	17382
NT2RP2004594	C-NT2RP2004594	17383	17384
NT2RP2004681	C-NT2RP2004681	17385	17386
NT2RP2004709	C-NT2RP2004709	17387	
NT2RP2004710	C-NT2RP2004710	17388	17389
NT2RP2004732	C-NT2RP2004732	17390	17391
NT2RP2004767	C-NT2RP2004767	17392	
NT2RP2004775	C-NT2RP2004775	17393	17394
NT2RP2004961	C-NT2RP2004961	17395	17396
NT2RP2004962	C-NT2RP2004962	17397	
NT2RP2004982	C-NT2RP2004982	17398	
NT2RP2005003	C-NT2RP2005003	17399	17400
NT2RP2005018	C-NT2RP2005018	17401	
NT2RP2005020	C-NT2RP2005020	17402	17403
NT2RP2005022	C-NT2RP2005022	17404	17405
NT2RP2005031	C-NT2RP2005031	17406	17407
NT2RP2005116	C-NT2RP2005116	17408	17409
NT2RP2005139	C-NT2RP2005139	17410	17411
NT2RP2005168	C-NT2RP2005168	17412	17413
NT2RP2005254	C-NT2RP2005254	17414	17415
NT2RP2005325	C-NT2RP2005325	17416	17417
NT2RP2005336	C-NT2RP2005336	17418	17419
NT2RP2005344	C-NT2RP2005344	17420	17421
NT2RP2005360	C-NT2RP2005360	17422	17423
NT2RP2005407	C-NT2RP2005407	17424	17425
NT2RP2005454	C-NT2RP2005454	17426	
NT2RP2005457	C-NT2RP2005457	17427	17428

(1000)

特開 2002-191363

1997

1998

NT2RP2005476	C-NT2RP2005476	17429	
NT2RP2005491	C-NT2RP2005491	17430	17431
NT2RP2005496	C-NT2RP2005496	17432	
NT2RP2005501	C-NT2RP2005501	17433	
NT2RP2005531	C-NT2RP2005531	17434	17435
NT2RP2005600	C-NT2RP2005600	17436	
NT2RP2005645	C-NT2RP2005645	17437	
NT2RP2005694	C-NT2RP2005694	17438	17439
NT2RP2005701	C-NT2RP2005701	17440	17441
NT2RP2005741	C-NT2RP2005741	17442	
NT2RP2005806	C-NT2RP2005806	17443	17444
NT2RP2005815	C-NT2RP2005815	17445	17446
NT2RP2005841	C-NT2RP2005841	17447	17448
NT2RP2005882	C-NT2RP2005882	17449	
NT2RP2005942	C-NT2RP2005942	17450	17451
NT2RP2006103	C-NT2RP2006103	17452	17453
NT2RP2006166	C-NT2RP2006166	17454	17455
NT2RP2006258	C-NT2RP2006258	17456	
NT2RP2006261	C-NT2RP2006261	17457	17458
NT2RP2006321	C-NT2RP2006321	17459	
NT2RP2006454	C-NT2RP2006454	17460	
NT2RP2006598	C-NT2RP2006598	17461	17462
NT2RP3000046	C-NT2RP3000046	17463	17464
NT2RP3000047	C-NT2RP3000047	17465	17466
NT2RP3000418	C-NT2RP3000418	17467	
NT2RP3000439	C-NT2RP3000439	17468	17469
NT2RP3000487	C-NT2RP3000487	17470	17471
NT2RP3000512	C-NT2RP3000512	17472	17473
NT2RP3000526	C-NT2RP3000526	17474	
NT2RP3000603	C-NT2RP3000603	17475	17476
NT2RP3000605	C-NT2RP3000605	17477	17478
NT2RP3000628	C-NT2RP3000628	17479	
NT2RP3000739	C-NT2RP3000739	17480	17481
NT2RP3000845	C-NT2RP3000845	17482	17483
NT2RP3000968	C-NT2RP3000968	17484	
NT2RP3001057	C-NT2RP3001057	17485	17486
NT2RP3001113	C-NT2RP3001113	17487	17488
NT2RP3001245	C-NT2RP3001245	17489	
NT2RP3001253	C-NT2RP3001253	17490	17491
NT2RP3001356	C-NT2RP3001356	17492	17493
NT2RP3001383	C-NT2RP3001383	17494	17495
NT2RP3001399	C-NT2RP3001399	17496	17497
NT2RP3001554	C-NT2RP3001554	17498	17499
NT2RP3001712	C-NT2RP3001712	17500	17501
NT2RP3001724	C-NT2RP3001724	17502	17503
NT2RP3001727	C-NT2RP3001727	17504	17505
NT2RP3001730	C-NT2RP3001730	17506	17507
NT2RP3001739	C-NT2RP3001739	17508	17509
NT2RP3001777	C-NT2RP3001777	17510	17511
NT2RP3001857	C-NT2RP3001857	17512	17513

(1001)

特開 2002-191363

1999

2000

NT2RP3001943	C-NT2RP3001943	17514	17515
NT2RP3001944	C-NT2RP3001944	17516	17517
NT2RP3002033	C-NT2RP3002033	17518	17519
NT2RP3002054	C-NT2RP3002054	17520	17521
NT2RP3002063	C-NT2RP3002063	17522	17523
NT2RP3002099	C-NT2RP3002099	17524	17525
NT2RP3002102	C-NT2RP3002102	17526	
NT2RP3002147	C-NT2RP3002147	17527	17528
NT2RP3002163	C-NT2RP3002163	17529	17530
NT2RP3002173	C-NT2RP3002173	17531	
NT2RP3002255	C-NT2RP3002255	17532	
NT2RP3002303	C-NT2RP3002303	17533	17534
NT2RP3002343	C-NT2RP3002343	17535	17536
NT2RP3002351	C-NT2RP3002351	17537	17538
NT2RP3002399	C-NT2RP3002399	17539	17540
NT2RP3002455	C-NT2RP3002455	17541	17542
NT2RP3002545	C-NT2RP3002545	17543	17544
NT2RP3002549	C-NT2RP3002549	17545	17546
NT2RP3002602	C-NT2RP3002602	17547	17548
NT2RP3002603	C-NT2RP3002603	17549	17550
NT2RP3002628	C-NT2RP3002628	17551	17552
NT2RP3002659	C-NT2RP3002659	17553	17554
NT2RP3002660	C-NT2RP3002660	17555	17556
NT2RP3002682	C-NT2RP3002682	17557	17558
NT2RP3002687	C-NT2RP3002687	17559	17560
NT2RP3002688	C-NT2RP3002688	17561	17562
NT2RP3002701	C-NT2RP3002701	17563	17564
NT2RP3002785	C-NT2RP3002785	17565	17566
NT2RP3002869	C-NT2RP3002869	17567	17568
NT2RP3002876	C-NT2RP3002876	17569	17570
NT2RP3002877	C-NT2RP3002877	17571	
NT2RP3002909	C-NT2RP3002909	17572	17573
NT2RP3002969	C-NT2RP3002969	17574	17575
NT2RP3002972	C-NT2RP3002972	17576	17577
NT2RP3003032	C-NT2RP3003032	17578	
NT2RP3003061	C-NT2RP3003061	17579	17580
NT2RP3003071	C-NT2RP3003071	17581	17582
NT2RP3003078	C-NT2RP3003078	17583	17584
NT2RP3003139	C-NT2RP3003139	17585	
NT2RP3003145	C-NT2RP3003145	17586	17587
NT2RP3003150	C-NT2RP3003150	17588	
NT2RP3003197	C-NT2RP3003197	17589	17590
NT2RP3003203	C-NT2RP3003203	17591	17592
NT2RP3003210	C-NT2RP3003210	17593	
NT2RP3003212	C-NT2RP3003212	17594	17595
NT2RP3003230	C-NT2RP3003230	17596	17597
NT2RP3003242	C-NT2RP3003242	17598	17599
NT2RP3003251	C-NT2RP3003251	17600	17601
NT2RP3003301	C-NT2RP3003301	17602	17603
NT2RP3003311	C-NT2RP3003311	17604	17605

(1002)

特開 2002-191363

2001

2002

NT2RP3003409	C-NT2RP3003409	17606	17607
NT2RP3003427	C-NT2RP3003427	17608	17609
NT2RP3003543	C-NT2RP3003543	17610	17611
NT2RP3003552	C-NT2RP3003552	17612	17613
NT2RP3003555	C-NT2RP3003555	17614	17615
NT2RP3003564	C-NT2RP3003564	17616	17617
NT2RP3003589	C-NT2RP3003589	17618	17619
NT2RP3003621	C-NT2RP3003621	17620	17621
NT2RP3003625	C-NT2RP3003625	17622	17623
NT2RP3003656	C-NT2RP3003656	17624	17625
NT2RP3003659	C-NT2RP3003659	17626	17627
NT2RP3003686	C-NT2RP3003686	17628	17629
NT2RP3003701	C-NT2RP3003701	17630	17631
NT2RP3003716	C-NT2RP3003716	17632	17633
NT2RP3003726	C-NT2RP3003726	17634	17635
NT2RP3003795	C-NT2RP3003795	17636	17637
NT2RP3003805	C-NT2RP3003805	17638	17639
NT2RP3003809	C-NT2RP3003809	17640	17641
NT2RP3003819	C-NT2RP3003819	17642	17643
NT2RP3003825	C-NT2RP3003825	17644	17645
NT2RP3003831	C-NT2RP3003831	17646	17647
NT2RP3003833	C-NT2RP3003833	17648	
NT2RP3003842	C-NT2RP3003842	17649	
NT2RP3003846	C-NT2RP3003846	17650	17651
NT2RP3003870	C-NT2RP3003870	17652	17653
NT2RP3003876	C-NT2RP3003876	17654	17655
NT2RP3003914	C-NT2RP3003914	17656	
NT2RP3003918	C-NT2RP3003918	17657	17658
NT2RP3003989	C-NT2RP3003989	17659	
NT2RP3004016	C-NT2RP3004016	17660	17661
NT2RP3004070	C-NT2RP3004070	17662	
NT2RP3004145	C-NT2RP3004145	17663	17664
NT2RP3004215	C-NT2RP3004215	17665	
NT2RP3004253	C-NT2RP3004253	17666	17667
NT2RP3004282	C-NT2RP3004282	17668	17669
NT2RP3004348	C-NT2RP3004348	17670	17671
NT2RP3004490	C-NT2RP3004490	17672	17673
NT2RP3004503	C-NT2RP3004503	17674	17675
NT2RP3004566	C-NT2RP3004566	17676	17677
NT2RP3004670	C-NT2RP3004670	17678	17679
NT2RP4000023	C-NT2RP4000023	17680	
NT2RP4000218	C-NT2RP4000218	17681	
NT2RP4000424	C-NT2RP4000424	17682	
NT2RP4001213	C-NT2RP4001213	17683	17684
NT2RP4001447	C-NT2RP4001447	17685	17686
NT2RP4001841	C-NT2RP4001841	17687	17688
NT2RP4001849	C-NT2RP4001849	17689	17690
NT2RP4002047	C-NT2RP4002047	17691	17692
NT2RP4002075	C-NT2RP4002075	17693	17694
NT2RP4002083	C-NT2RP4002083	17695	17696

(1003)

特開 2002-191363

2003

2004

OVARC1000001	C-OVARC1000001	17697	17698
OVARC1000008	C-OVARC1000008	17699	17700
OVARC1000017	C-OVARC1000017	17701	17702
OVARC1000058	C-OVARC1000058	17703	17704
OVARC1000068	C-OVARC1000068	17705	
OVARC1000071	C-OVARC1000071	17706	17707
OVARC1000085	C-OVARC1000085	17708	17709
OVARC1000109	C-OVARC1000109	17710	17711
OVARC1000114	C-OVARC1000114	17712	17713
OVARC1000145	C-OVARC1000145	17714	17715
OVARC1000240	C-OVARC1000240	17716	
OVARC1000302	C-OVARC1000302	17717	
OVARC1000408	C-OVARC1000408	17718	17719
OVARC1000414	C-OVARC1000414	17720	
OVARC1000440	C-OVARC1000440	17721	
OVARC1000442	C-OVARC1000442	17722	
OVARC1000496	C-OVARC1000496	17723	
OVARC1000556	C-OVARC1000556	17724	17725
OVARC1000557	C-OVARC1000557	17726	17727
OVARC1000578	C-OVARC1000578	17728	
OVARC1000622	C-OVARC1000622	17729	
OVARC1000679	C-OVARC1000679	17730	17731
OVARC1000681	C-OVARC1000681	17732	
OVARC1000700	C-OVARC1000700	17733	17734
OVARC1000724	C-OVARC1000724	17735	17736
OVARC1000751	C-OVARC1000751	17737	17738
OVARC1000800	C-OVARC1000800	17739	
OVARC1000885	C-OVARC1000885	17740	17741
OVARC1000936	C-OVARC1000936	17742	17743
OVARC1000937	C-OVARC1000937	17744	17745
OVARC1000960	C-OVARC1000960	17746	
OVARC1000971	C-OVARC1000971	17747	
OVARC1000999	C-OVARC1000999	17748	17749
OVARC1001000	C-OVARC1001000	17750	
OVARC1001029	C-OVARC1001029	17751	17752
OVARC1001040	C-OVARC1001040	17753	
OVARC1001051	C-OVARC1001051	17754	17755
OVARC1001113	C-OVARC1001113	17756	17757
OVARC1001118	C-OVARC1001118	17758	
OVARC1001129	C-OVARC1001129	17759	17760
OVARC1001169	C-OVARC1001169	17761	
OVARC1001240	C-OVARC1001240	17762	
OVARC1001261	C-OVARC1001261	17763	
OVARC1001339	C-OVARC1001339	17764	
OVARC1001342	C-OVARC1001342	17765	17766
OVARC1001357	C-OVARC1001357	17767	
OVARC1001442	C-OVARC1001442	17768	
OVARC1001611	C-OVARC1001611	17769	
OVARC1001813	C-OVARC1001813	17770	
OVARC1002112	C-OVARC1002112	17771	17772

(1004)

特開 2002-191363

2005

2006

OVARC1002143	C-OVARC1002143	17773	
OVARC1002165	C-OVARC1002165	17774	17775
OVARC1002182	C-OVARC1002182	17776	17777
PLACE1000014	C-PLACE1000014	17778	17779
PLACE1000078	C-PLACE1000078	17780	
PLACE1000492	C-PLACE1000492	17781	17782
PLACE1000793	C-PLACE1000793	17783	17784
PLACE1000814	C-PLACE1000814	17785	17786
PLACE1000979	C-PLACE1000979	17787	17788
PLACE1001007	C-PLACE1001007	17789	
PLACE1001054	C-PLACE1001054	17790	17791
PLACE1001088	C-PLACE1001088	17792	17793
PLACE1001136	C-PLACE1001136	17794	
PLACE1001241	C-PLACE1001241	17795	17796
PLACE1001377	C-PLACE1001377	17797	17798
PLACE1001395	C-PLACE1001395	17799	17800
PLACE1001740	C-PLACE1001740	17801	
PLACE1001746	C-PLACE1001746	17802	
PLACE1001983	C-PLACE1001983	17803	17804
PLACE1002066	C-PLACE1002066	17805	
PLACE1002115	C-PLACE1002115	17806	
PLACE1002213	C-PLACE1002213	17807	17808
PLACE1002342	C-PLACE1002342	17809	17810
PLACE1002450	C-PLACE1002450	17811	17812
PLACE1002474	C-PLACE1002474	17813	17814
PLACE1002499	C-PLACE1002499	17815	17816
PLACE1002578	C-PLACE1002578	17817	
PLACE1002714	C-PLACE1002714	17818	17819
PLACE1002772	C-PLACE1002772	17820	17821
PLACE1002775	C-PLACE1002775	17822	17823
PLACE1002834	C-PLACE1002834	17824	17825
PLACE1002993	C-PLACE1002993	17826	17827
PLACE1003030	C-PLACE1003030	17828	17829
PLACE1003205	C-PLACE1003205	17830	
PLACE1003249	C-PLACE1003249	17831	
PLACE1003493	C-PLACE1003493	17832	17833
PLACE1003553	C-PLACE1003553	17834	17835
PLACE1003592	C-PLACE1003592	17836	
PLACE1003596	C-PLACE1003596	17837	17838
PLACE1003669	C-PLACE1003669	17839	17840
PLACE1003709	C-PLACE1003709	17841	17842
PLACE1003870	C-PLACE1003870	17843	17844
PLACE1003885	C-PLACE1003885	17845	17846
PLACE1003892	C-PLACE1003892	17847	
PLACE1003900	C-PLACE1003900	17848	
PLACE1004336	C-PLACE1004336	17849	
PLACE1004384	C-PLACE1004384	17850	
PLACE1004425	C-PLACE1004425	17851	
PLACE1004471	C-PLACE1004471	17852	
PLACE1004506	C-PLACE1004506	17853	17854

(1005)

特開 2002-191363

2007

2008

PLACE1004518	C-PLACE1004518	17855	
PLACE1004550	C-PLACE1004550	17856	17857
PLACE1004681	C-PLACE1004681	17858	
PLACE1004693	C-PLACE1004693	17859	
PLACE1004716	C-PLACE1004716	17860	
PLACE1004815	C-PLACE1004815	17861	17862
PLACE1004836	C-PLACE1004836	17863	17864
PLACE1004838	C-PLACE1004838	17865	17866
PLACE1004840	C-PLACE1004840	17867	17868
PLACE1004900	C-PLACE1004900	17869	
PLACE1004985	C-PLACE1004985	17870	
PLACE1005085	C-PLACE1005085	17871	
PLACE1005086	C-PLACE1005086	17872	17873
PLACE1005108	C-PLACE1005108	17874	
PLACE1005146	C-PLACE1005146	17875	
PLACE1005409	C-PLACE1005409	17876	
PLACE1005453	C-PLACE1005453	17877	
PLACE1005477	C-PLACE1005477	17878	17879
PLACE1005557	C-PLACE1005557	17880	
PLACE1005595	C-PLACE1005595	17881	17882
PLACE1005603	C-PLACE1005603	17883	17884
PLACE1005639	C-PLACE1005639	17885	
PLACE1005727	C-PLACE1005727	17886	17887
PLACE1005799	C-PLACE1005799	17888	17889
PLACE1005813	C-PLACE1005813	17890	17891
PLACE1005884	C-PLACE1005884	17892	17893
PLACE1005968	C-PLACE1005968	17894	
PLACE1006002	C-PLACE1006002	17895	
PLACE1006003	C-PLACE1006003	17896	17897
PLACE1006017	C-PLACE1006017	17898	17899
PLACE1006037	C-PLACE1006037	17900	17901
PLACE1006076	C-PLACE1006076	17902	
PLACE1006143	C-PLACE1006143	17903	
PLACE1006248	C-PLACE1006248	17904	17905
PLACE1006288	C-PLACE1006288	17906	
PLACE1006318	C-PLACE1006318	17907	17908
PLACE1006368	C-PLACE1006368	17909	17910
PLACE1006371	C-PLACE1006371	17911	
PLACE1006469	C-PLACE1006469	17912	17913
PLACE1006506	C-PLACE1006506	17914	17915
PLACE1006521	C-PLACE1006521	17916	
PLACE1006534	C-PLACE1006534	17917	
PLACE1006617	C-PLACE1006617	17918	
PLACE1006640	C-PLACE1006640	17919	
PLACE1006754	C-PLACE1006754	17920	17921
PLACE1006760	C-PLACE1006760	17922	17923
PLACE1006779	C-PLACE1006779	17924	
PLACE1006805	C-PLACE1006805	17925	
PLACE1006815	C-PLACE1006815	17926	17927
PLACE1006867	C-PLACE1006867	17928	

(1006)

特開 2002-191363

2009

2010

PLACE1007045	C-PLACE1007045	17929	17930
PLACE1007097	C-PLACE1007097	17931	
PLACE1007111	C-PLACE1007111	17932	17933
PLACE1007112	C-PLACE1007112	17934	17935
PLACE1007140	C-PLACE1007140	17936	17937
PLACE1007218	C-PLACE1007218	17938	17939
PLACE1007454	C-PLACE1007454	17940	
PLACE1007478	C-PLACE1007478	17941	
PLACE1007677	C-PLACE1007677	17942	
PLACE1007705	C-PLACE1007705	17943	17944
PLACE1007737	C-PLACE1007737	17945	
PLACE1007743	C-PLACE1007743	17946	
PLACE1007852	C-PLACE1007852	17947	17948
PLACE1007877	C-PLACE1007877	17949	17950
PLACE1008045	C-PLACE1008045	17951	
PLACE1008080	C-PLACE1008080	17952	17953
PLACE1008111	C-PLACE1008111	17954	17955
PLACE1008201	C-PLACE1008201	17956	17957
PLACE1008231	C-PLACE1008231	17958	
PLACE1008244	C-PLACE1008244	17959	17960
PLACE1008330	C-PLACE1008330	17961	
PLACE1008331	C-PLACE1008331	17962	17963
PLACE1008369	C-PLACE1008369	17964	
PLACE1008392	C-PLACE1008392	17965	
PLACE1008405	C-PLACE1008405	17966	
PLACE1008424	C-PLACE1008424	17967	17968
PLACE1008584	C-PLACE1008584	17969	17970
PLACE1008625	C-PLACE1008625	17971	17972
PLACE1008630	C-PLACE1008630	17973	
PLACE1008643	C-PLACE1008643	17974	17975
PLACE1008715	C-PLACE1008715	17976	17977
PLACE1008748	C-PLACE1008748	17978	
PLACE1008757	C-PLACE1008757	17979	
PLACE1008798	C-PLACE1008798	17980	
PLACE1008851	C-PLACE1008851	17981	
PLACE1008947	C-PLACE1008947	17982	17983
PLACE1009039	C-PLACE1009039	17984	
PLACE1009048	C-PLACE1009048	17985	
PLACE1009050	C-PLACE1009050	17986	
PLACE1009113	C-PLACE1009113	17987	17988
PLACE1009150	C-PLACE1009150	17989	
PLACE1009200	C-PLACE1009200	17990	
PLACE1009246	C-PLACE1009246	17991	17992
PLACE1009298	C-PLACE1009298	17993	17994
PLACE1009308	C-PLACE1009308	17995	17996
PLACE1009398	C-PLACE1009398	17997	17998
PLACE1009410	C-PLACE1009410	17999	
PLACE1009477	C-PLACE1009477	18000	
PLACE1009493	C-PLACE1009493	18001	18002
PLACE1009539	C-PLACE1009539	18003	

(1007)

特開 2002-191363

2011

2012

PLACE1009595	C-PLACE1009595	18004	
PLACE1009637	C-PLACE1009637	18005	
PLACE1009639	C-PLACE1009639	18006	18007
PLACE1009798	C-PLACE1009798	18008	18009
PLACE1009861	C-PLACE1009861	18010	18011
PLACE1009888	C-PLACE1009888	18012	
PLACE1009925	C-PLACE1009925	18013	18014
PLACE1009947	C-PLACE1009947	18015	
PLACE1010069	C-PLACE1010069	18016	
PLACE1010089	C-PLACE1010089	18017	18018
PLACE1010231	C-PLACE1010231	18019	18020
PLACE1010270	C-PLACE1010270	18021	
PLACE1010562	C-PLACE1010562	18022	18023
PLACE1010579	C-PLACE1010579	18024	18025
PLACE1010624	C-PLACE1010624	18026	18027
PLACE1010628	C-PLACE1010628	18028	18029
PLACE1010662	C-PLACE1010662	18030	18031
PLACE1010702	C-PLACE1010702	18032	18033
PLACE1010761	C-PLACE1010761	18034	18035
PLACE1010802	C-PLACE1010802	18036	18037
PLACE1010833	C-PLACE1010833	18038	18039
PLACE1010896	C-PLACE1010896	18040	18041
PLACE1010916	C-PLACE1010916	18042	
PLACE1010947	C-PLACE1010947	18043	
PLACE1010965	C-PLACE1010965	18044	
PLACE1011032	C-PLACE1011032	18045	
PLACE1011041	C-PLACE1011041	18046	18047
PLACE1011056	C-PLACE1011056	18048	18049
PLACE1011090	C-PLACE1011090	18050	18051
PLACE1011160	C-PLACE1011160	18052	18053
PLACE1011214	C-PLACE1011214	18054	18055
PLACE1011229	C-PLACE1011229	18056	18057
PLACE1011273	C-PLACE1011273	18058	
PLACE1011291	C-PLACE1011291	18059	18060
PLACE1011310	C-PLACE1011310	18061	18062
PLACE1011371	C-PLACE1011371	18063	18064
PLACE1011503	C-PLACE1011503	18065	
PLACE1011635	C-PLACE1011635	18066	18067
PLACE1011646	C-PLACE1011646	18068	18069
PLACE1011650	C-PLACE1011650	18070	
PLACE1011675	C-PLACE1011675	18071	
PLACE1011725	C-PLACE1011725	18072	18073
PLACE1011749	C-PLACE1011749	18074	18075
PLACE1011922	C-PLACE1011922	18076	18077
PLACE1012031	C-PLACE1012031	18078	18079
PLACE2000006	C-PLACE2000006	18080	18081
PLACE2000007	C-PLACE2000007	18082	18083
PLACE2000034	C-PLACE2000034	18084	18085
PLACE2000039	C-PLACE2000039	18086	18087
PLACE2000061	C-PLACE2000061	18088	18089

(1008)

特開 2002-191363

2013

2014

PLACE2000072	C-PLACE2000072	18090	18091
PLACE2000097	C-PLACE2000097	18092	18093
PLACE2000103	C-PLACE2000103	18094	
PLACE2000115	C-PLACE2000115	18095	
PLACE2000124	C-PLACE2000124	18096	18097
PLACE2000140	C-PLACE2000140	18098	
PLACE2000164	C-PLACE2000164	18099	18100
PLACE2000176	C-PLACE2000176	18101	18102
PLACE2000223	C-PLACE2000223	18103	
PLACE2000235	C-PLACE2000235	18104	
PLACE2000274	C-PLACE2000274	18105	18106
PLACE2000302	C-PLACE2000302	18107	18108
PLACE2000347	C-PLACE2000347	18109	
PLACE2000359	C-PLACE2000359	18110	18111
PLACE2000371	C-PLACE2000371	18112	18113
PLACE2000379	C-PLACE2000379	18114	18115
PLACE2000399	C-PLACE2000399	18116	
PLACE2000404	C-PLACE2000404	18117	18118
PLACE2000450	C-PLACE2000450	18119	
PLACE2000455	C-PLACE2000455	18120	
PLACE3000059	C-PLACE3000059	18121	18122
PLACE3000070	C-PLACE3000070	18123	
PLACE3000119	C-PLACE3000119	18124	
PLACE3000121	C-PLACE3000121	18125	18126
PLACE3000136	C-PLACE3000136	18127	18128
PLACE3000147	C-PLACE3000147	18129	18130
PLACE3000148	C-PLACE3000148	18131	18132
PLACE3000155	C-PLACE3000155	18133	18134
PLACE3000160	C-PLACE3000160	18135	18136
PLACE3000169	C-PLACE3000169	18137	18138
PLACE3000194	C-PLACE3000194	18139	
PLACE3000199	C-PLACE3000199	18140	18141
PLACE3000218	C-PLACE3000218	18142	18143
PLACE3000230	C-PLACE3000230	18144	18145
PLACE3000244	C-PLACE3000244	18146	18147
PLACE3000254	C-PLACE3000254	18148	18149
PLACE3000276	C-PLACE3000276	18150	
PLACE3000310	C-PLACE3000310	18151	
PLACE3000320	C-PLACE3000320	18152	
PLACE3000331	C-PLACE3000331	18153	
PLACE3000339	C-PLACE3000339	18154	18155
PLACE3000352	C-PLACE3000352	18156	
PLACE3000353	C-PLACE3000353	18157	18158
PLACE3000362	C-PLACE3000362	18159	18160
PLACE3000365	C-PLACE3000365	18161	
PLACE3000388	C-PLACE3000388	18162	
PLACE3000413	C-PLACE3000413	18163	18164
PLACE3000425	C-PLACE3000425	18165	
PLACE4000009	C-PLACE4000009	18166	18167
PLACE4000014	C-PLACE4000014	18168	18169

(1009)

特開 2002-191363

2015

2016

PLACE4000052	C-PLACE4000052	18170	18171
PLACE4000089	C-PLACE4000089	18172	18173
PLACE4000128	C-PLACE4000128	18174	18175
PLACE4000129	C-PLACE4000129	18176	18177
PLACE4000147	C-PLACE4000147	18178	18179
PLACE4000192	C-PLACE4000192	18180	18181
PLACE4000211	C-PLACE4000211	18182	18183
PLACE4000222	C-PLACE4000222	18184	18185
PLACE4000269	C-PLACE4000269	18186	18187
PLACE4000270	C-PLACE4000270	18188	18189
PLACE4000300	C-PLACE4000300	18190	18191
PLACE4000387	C-PLACE4000387	18192	18193
PLACE4000392	C-PLACE4000392	18194	18195
PLACE4000431	C-PLACE4000431	18196	18197
PLACE4000450	C-PLACE4000450	18198	18199
PLACE4000465	C-PLACE4000465	18200	
PLACE4000489	C-PLACE4000489	18201	18202
PLACE4000654	C-PLACE4000654	18203	18204
SKNMC1000011	C-SKNMC1000011	18205	18206
SKNMC1000046	C-SKNMC1000046	18207	18208
SKNMC1000050	C-SKNMC1000050	18209	
THYRO1000034	C-THYRO1000034	18210	18211
THYRO1000070	C-THYRO1000070	18212	18213
THYRO1000072	C-THYRO1000072	18214	18215
THYRO1000092	C-THYRO1000092	18216	18217
THYRO1000121	C-THYRO1000121	18218	18219
THYRO1000124	C-THYRO1000124	18220	18221
THYRO1000197	C-THYRO1000197	18222	18223
THYRO1000199	C-THYRO1000199	18224	18225
THYRO1000206	C-THYRO1000206	18226	
THYRO1000242	C-THYRO1000242	18227	18228
THYRO1000253	C-THYRO1000253	18229	18230
THYRO1000270	C-THYRO1000270	18231	18232
THYRO1000288	C-THYRO1000288	18233	18234
THYRO1000320	C-THYRO1000320	18235	
THYRO1000358	C-THYRO1000358	18236	18237
THYRO1000368	C-THYRO1000368	18238	18239
THYRO1000381	C-THYRO1000381	18240	18241
THYRO1000387	C-THYRO1000387	18242	
THYRO1000394	C-THYRO1000394	18243	18244
THYRO1000395	C-THYRO1000395	18245	18246
THYRO1000401	C-THYRO1000401	18247	18248
THYRO1000488	C-THYRO1000488	18249	18250
THYRO1000501	C-THYRO1000501	18251	18252
THYRO1000558	C-THYRO1000558	18253	
THYRO1000570	C-THYRO1000570	18254	18255
THYRO1000605	C-THYRO1000605	18256	18257
THYRO1000625	C-THYRO1000625	18258	
THYRO1000637	C-THYRO1000637	18259	
THYRO1000676	C-THYRO1000676	18260	

(1010)

特開 2002-191363

2017

2018

THYRO1000684	C-THYRO1000684	18261	18262
THYRO1000712	C-THYRO1000712	18263	
THYRO1000805	C-THYRO1000805	18264	
THYRO1000815	C-THYRO1000815	18265	
THYRO1000855	C-THYRO1000855	18266	18267
THYRO1000934	C-THYRO1000934	18268	18269
THYRO1000988	C-THYRO1000988	18270	18271
THYRO1001033	C-THYRO1001033	18272	18273
THYRO1001120	C-THYRO1001120	18274	18275
THYRO1001204	C-THYRO1001204	18276	18277
THYRO1001262	C-THYRO1001262	18278	18279
THYRO1001271	C-THYRO1001271	18280	18281
THYRO1001287	C-THYRO1001287	18282	18283
THYRO1001313	C-THYRO1001313	18284	18285
THYRO1001347	C-THYRO1001347	18286	18287
THYRO1001363	C-THYRO1001363	18288	
THYRO1001374	C-THYRO1001374	18289	18290
THYRO1001403	C-THYRO1001403	18291	18292
THYRO1001405	C-THYRO1001405	18293	18294
THYRO1001406	C-THYRO1001406	18295	18296
THYRO1001426	C-THYRO1001426	18297	
THYRO1001458	C-THYRO1001458	18298	18299
THYRO1001480	C-THYRO1001480	18300	18301
THYRO1001487	C-THYRO1001487	18302	
THYRO1001584	C-THYRO1001584	18303	18304
THYRO1001661	C-THYRO1001661	18305	18306
THYRO1001746	C-THYRO1001746	18307	
THYRO1001772	C-THYRO1001772	18308	
THYRO1001854	C-THYRO1001854	18309	
Y79AA1000013	C-Y79AA1000013	18310	18311
Y79AA1000033	C-Y79AA1000033	18312	18313
Y79AA1000231	C-Y79AA1000231	18314	18315
Y79AA1000342	C-Y79AA1000342	18316	18317
Y79AA1000349	C-Y79AA1000349	18318	18319
Y79AA1000410	C-Y79AA1000410	18320	
Y79AA1000539	C-Y79AA1000539	18321	18322
Y79AA1000589	C-Y79AA1000589	18323	18324
Y79AA1000802	C-Y79AA1000802	18325	18326
Y79AA1000827	C-Y79AA1000827	18327	18328
Y79AA1000966	C-Y79AA1000966	18329	18330
Y79AA1000969	C-Y79AA1000969	18331	
Y79AA1000985	C-Y79AA1000985	18332	18333
Y79AA1001061	C-Y79AA1001061	18334	
Y79AA1001068	C-Y79AA1001068	18335	18336
Y79AA1001216	C-Y79AA1001216	18337	18338
Y79AA1001299	C-Y79AA1001299	18339	18340
Y79AA1001511	C-Y79AA1001511	18341	18342
Y79AA1001594	C-Y79AA1001594	18343	18344
Y79AA1001692	C-Y79AA1001692	18345	18346
Y79AA1001866	C-Y79AA1001866	18347	18348

(1011)

特開 2002-191363

2019

2020

Y79AA1001874	C-Y79AA1001874	18349	18350
Y79AA1002139	C-Y79AA1002139	18351	18352
Y79AA1002210	C-Y79AA1002210	18353	18354
Y79AA1002211	C-Y79AA1002211	18355	18356
Y79AA1002220	C-Y79AA1002220	18357	
Y79AA1002234	C-Y79AA1002234	18358	18359
Y79AA1002258	C-Y79AA1002258	18360	18361
Y79AA1002361	C-Y79AA1002361	18362	18363
Y79AA1002399	C-Y79AA1002399	18364	18365
Y79AA1002416	C-Y79AA1002416	18366	18367
Y79AA1002431	C-Y79AA1002431	18368	18369
Y79AA1002482	C-Y79AA1002482	18370	18371
Y79AA1002487	C-Y79AA1002487	18372	18373
HEMBA1000459	C-HEMBA1000459	18374	
HEMBA1000505	C-HEMBA1000505	18375	18376
HEMBA1002503	C-HEMBA1002503	18377	
HEMBA1002508	C-HEMBA1002508	18378	
HEMBA1002513	C-HEMBA1002513	18379	18380
HEMBA1003480	C-HEMBA1003480	18381	18382
HEMBA1003538	C-HEMBA1003538	18383	18384
HEMBA1003645	C-HEMBA1003645	18385	18386
HEMBA1003646	C-HEMBA1003646	18387	18388
HEMBA1003667	C-HEMBA1003667	18389	
HEMBA1003679	C-HEMBA1003679	18390	18391
HEMBA1003827	C-HEMBA1003827	18392	18393
HEMBA1003838	C-HEMBA1003838	18394	
HEMBA1004055	C-HEMBA1004055	18395	18396
HEMBA1004056	C-HEMBA1004056	18397	
HEMBA1004086	C-HEMBA1004086	18398	
HEMBA1004335	C-HEMBA1004335	18399	18400
HEMBA1004353	C-HEMBA1004353	18401	
HEMBA1004479	C-HEMBA1004479	18402	18403
HEMBA1004499	C-HEMBA1004499	18404	18405
HEMBA1004507	C-HEMBA1004507	18406	18407
HEMBA1004638	C-HEMBA1004638	18408	18409
HEMBA1004669	C-HEMBA1004669	18410	18411
HEMBA1004709	C-HEMBA1004709	18412	
HEMBA1004860	C-HEMBA1004860	18413	
HEMBA1005206	C-HEMBA1005206	18414	18415
HEMBA1005472	C-HEMBA1005472	18416	18417
HEMBA1005513	C-HEMBA1005513	18418	18419
HEMBA1005572	C-HEMBA1005572	18420	
HEMBA1005780	C-HEMBA1005780	18421	
HEMBA1005990	C-HEMBA1005990	18422	18423
HEMBA1006038	C-HEMBA1006038	18424	18425
HEMBA1006124	C-HEMBA1006124	18426	
HEMBA1006461	C-HEMBA1006461	18427	
HEMBA1006521	C-HEMBA1006521	18428	18429
HEMBA1006617	C-HEMBA1006617	18430	18431
HEMBA1006779	C-HEMBA1006779	18432	

(1012)

特開 2002-191363

2021

2022

HEMBA1006812	C-HEMBA1006812	18433	18434
HEMBA1006914	C-HEMBA1006914	18435	18436
HEMBA1007174	C-HEMBA1007174	18437	18438
HEMBA1000240	C-HEMBA1000240	18439	18440
HEMBA1000264	C-HEMBA1000264	18441	18442
HEMBA1000335	C-HEMBA1000335	18443	
HEMBA1000554	C-HEMBA1000554	18444	18445
HEMBA1000573	C-HEMBA1000573	18446	
HEMBA1000749	C-HEMBA1000749	18447	18448
HEMBA1000774	C-HEMBA1000774	18449	18450
HEMBA1000835	C-HEMBA1000835	18451	18452
HEMBA1001197	C-HEMBA1001197	18453	
HEMBA1001315	C-HEMBA1001315	18454	18455
HEMBA1001482	C-HEMBA1001482	18456	18457
HEMBA1001500	C-HEMBA1001500	18458	18459
HEMBA1001562	C-HEMBA1001562	18460	18461
HEMBA1001630	C-HEMBA1001630	18462	18463
HEMBA1001665	C-HEMBA1001665	18464	
HEMBA1001684	C-HEMBA1001684	18465	18466
HEMBA1001812	C-HEMBA1001812	18467	
HEMBA1001834	C-HEMBA1001834	18468	18469
HEMBA1001869	C-HEMBA1001869	18470	18471
HEMBA1001871	C-HEMBA1001871	18472	18473
HEMBA1001872	C-HEMBA1001872	18474	18475
HEMBA1001905	C-HEMBA1001905	18476	18477
HEMBA1001908	C-HEMBA1001908	18478	18479
HEMBA1001915	C-HEMBA1001915	18480	18481
HEMBA1001925	C-HEMBA1001925	18482	18483
HEMBA1002044	C-HEMBA1002044	18484	18485
HEMBA1002134	C-HEMBA1002134	18486	18487
HEMBA1002152	C-HEMBA1002152	18488	
HEMBA1002300	C-HEMBA1002300	18489	18490
HEMBA1002381	C-HEMBA1002381	18491	18492
HEMBA1002383	C-HEMBA1002383	18493	18494
HEMBA1002534	C-HEMBA1002534	18495	
MAMMA1000143	C-MAMMA1000143	18496	
MAMMA1000183	C-MAMMA1000183	18497	18498
MAMMA1000227	C-MAMMA1000227	18499	
MAMMA1000257	C-MAMMA1000257	18500	
MAMMA1000264	C-MAMMA1000264	18501	
MAMMA1000279	C-MAMMA1000279	18502	
MAMMA1000372	C-MAMMA1000372	18503	
MAMMA1000752	C-MAMMA1000752	18504	
MAMMA1000760	C-MAMMA1000760	18505	18506
MAMMA1000778	C-MAMMA1000778	18507	18508
MAMMA1000855	C-MAMMA1000855	18509	18510
MAMMA1000859	C-MAMMA1000859	18511	18512
MAMMA1000897	C-MAMMA1000897	18513	18514
MAMMA1001073	C-MAMMA1001073	18515	18516
MAMMA1001080	C-MAMMA1001080	18517	18518

(1013)

特開 2002-191363

2023

2024

MAMMA1001198	C-MAMMA1001198	18519	18520
MAMMA1001202	C-MAMMA1001202	18521	18522
MAMMA1001222	C-MAMMA1001222	18523	18524
MAMMA1001252	C-MAMMA1001252	18525	18526
MAMMA1001296	C-MAMMA1001296	18527	
MAMMA1001502	C-MAMMA1001502	18528	18529
MAMMA1001630	C-MAMMA1001630	18530	
MAMMA1001633	C-MAMMA1001633	18531	18532
MAMMA1001683	C-MAMMA1001683	18533	
MAMMA1001715	C-MAMMA1001715	18534	
MAMMA1001730	C-MAMMA1001730	18535	18536
MAMMA1001760	C-MAMMA1001760	18537	18538
MAMMA1001769	C-MAMMA1001769	18539	
MAMMA1001785	C-MAMMA1001785	18540	18541
MAMMA1001848	C-MAMMA1001848	18542	
MAMMA1001874	C-MAMMA1001874	18543	
MAMMA1001956	C-MAMMA1001956	18544	
MAMMA1002009	C-MAMMA1002009	18545	18546
MAMMA1002033	C-MAMMA1002033	18547	
MAMMA1002155	C-MAMMA1002155	18548	
MAMMA1002498	C-MAMMA1002498	18549	18550
MAMMA1002571	C-MAMMA1002571	18551	18552
MAMMA1002573	C-MAMMA1002573	18553	18554
MAMMA1002590	C-MAMMA1002590	18555	18556
MAMMA1002617	C-MAMMA1002617	18557	18558
MAMMA1002618	C-MAMMA1002618	18559	
MAMMA1002636	C-MAMMA1002636	18560	
MAMMA1002646	C-MAMMA1002646	18561	18562
MAMMA1002665	C-MAMMA1002665	18563	
MAMMA1002708	C-MAMMA1002708	18564	
MAMMA1002728	C-MAMMA1002728	18565	
MAMMA1002744	C-MAMMA1002744	18566	
MAMMA1002764	C-MAMMA1002764	18567	18568
MAMMA1002765	C-MAMMA1002765	18569	
MAMMA1002830	C-MAMMA1002830	18570	
MAMMA1002844	C-MAMMA1002844	18571	18572
MAMMA1002858	C-MAMMA1002858	18573	18574
MAMMA1002880	C-MAMMA1002880	18575	18576
MAMMA1002892	C-MAMMA1002892	18577	
MAMMA1002909	C-MAMMA1002909	18578	
MAMMA1002941	C-MAMMA1002941	18579	18580
MAMMA1002947	C-MAMMA1002947	18581	18582
MAMMA1002973	C-MAMMA1002973	18583	
MAMMA1002987	C-MAMMA1002987	18584	
MAMMA1003003	C-MAMMA1003003	18585	
MAMMA1003026	C-MAMMA1003026	18586	18587
MAMMA1003031	C-MAMMA1003031	18588	
MAMMA1003089	C-MAMMA1003089	18589	18590
NT2RM1000092	C-NT2RM1000092	18591	18592
NT2RM1000272	C-NT2RM1000272	18593	18594

(1014)

特開 2002-191363

2025

2026

NT2RM1000341	C-NT2RM1000341	18595	18596
NT2RM1000539	C-NT2RM1000539	18597	18598
NT2RM1000553	C-NT2RM1000553	18599	18600
NT2RM1000623	C-NT2RM1000623	18601	18602
NT2RM1000702	C-NT2RM1000702	18603	18604
NT2RM1000833	C-NT2RM1000833	18605	18606
NT2RM1000883	C-NT2RM1000883	18607	18608
NT2RM1001082	C-NT2RM1001082	18609	
NT2RM1001112	C-NT2RM1001112	18610	18611
NT2RM2001105	C-NT2RM2001105	18612	18613
NT2RM2001360	C-NT2RM2001360	18614	18615
NT2RM2001797	C-NT2RM2001797	18616	18617
NT2RM2001803	C-NT2RM2001803	18618	18619
NT2RM4002504	C-NT2RM4002504	18620	
NT2RP1000409	C-NT2RP1000409	18621	
NT2RP1000460	C-NT2RP1000460	18622	18623
NT2RP1000746	C-NT2RP1000746	18624	18625
NT2RP1000796	C-NT2RP1000796	18626	18627
NT2RP1001013	C-NT2RP1001013	18628	18629
NT2RP2001214	C-NT2RP2001214	18630	18631
NT2RP2001233	C-NT2RP2001233	18632	18633
NT2RP2002056	C-NT2RP2002056	18634	18635
NT2RP2002105	C-NT2RP2002105	18636	18637
NT2RP2002333	C-NT2RP2002333	18638	18639
NT2RP2002677	C-NT2RP2002677	18640	
NT2RP2002755	C-NT2RP2002755	18641	
NT2RP2002843	C-NT2RP2002843	18642	
NT2RP2003101	C-NT2RP2003101	18643	18644
NT2RP2003668	C-NT2RP2003668	18645	18646
NT2RP2003799	C-NT2RP2003799	18647	
NT2RP2004095	C-NT2RP2004095	18648	18649
NT2RP2004300	C-NT2RP2004300	18650	18651
NT2RP2004675	C-NT2RP2004675	18652	18653
NT2RP2004920	C-NT2RP2004920	18654	18655
NT2RP2005144	C-NT2RP2005144	18656	18657
NT2RP2005719	C-NT2RP2005719	18658	
NT2RP2005726	C-NT2RP2005726	18659	18660
NT2RP2005776	C-NT2RP2005776	18661	18662
NT2RP2005980	C-NT2RP2005980	18663	
NT2RP2006184	C-NT2RP2006184	18664	18665
NT2RP2006534	C-NT2RP2006534	18666	
NT2RP2006554	C-NT2RP2006554	18667	18668
NT2RP3000584	C-NT2RP3000584	18669	
NT2RP3001115	C-NT2RP3001115	18670	
NT2RP3001723	C-NT2RP3001723	18671	18672
NT2RP3001938	C-NT2RP3001938	18673	18674
NT2RP3002330	C-NT2RP3002330	18675	18676
NT2RP3002402	C-NT2RP3002402	18677	18678
NT2RP3002484	C-NT2RP3002484	18679	18680
NT2RP3002512	C-NT2RP3002512	18681	18682

(1015)

特開 2002-191363

2027

2028

NT2RP3002713	C-NT2RP3002713	18683	
NT2RP3002799	C-NT2RP3002799	18684	
NT2RP3002810	C-NT2RP3002810	18685	18686
NT2RP3002818	C-NT2RP3002818	18687	18688
NT2RP3002955	C-NT2RP3002955	18689	
NT2RP3002985	C-NT2RP3002985	18690	18691
NT2RP3003059	C-NT2RP3003059	18692	18693
NT2RP3003121	C-NT2RP3003121	18694	18695
NT2RP3003133	C-NT2RP3003133	18696	18697
NT2RP3003155	C-NT2RP3003155	18698	18699
NT2RP3003157	C-NT2RP3003157	18700	18701
NT2RP3003185	C-NT2RP3003185	18702	18703
NT2RP3003264	C-NT2RP3003264	18704	18705
NT2RP3003346	C-NT2RP3003346	18706	
NT2RP3003403	C-NT2RP3003403	18707	
NT2RP3003411	C-NT2RP3003411	18708	18709
NT2RP3003500	C-NT2RP3003500	18710	18711
NT2RP3003572	C-NT2RP3003572	18712	18713
NT2RP3003576	C-NT2RP3003576	18714	18715
NT2RP3003665	C-NT2RP3003665	18716	18717
NT2RP3003672	C-NT2RP3003672	18718	
NT2RP3003680	C-NT2RP3003680	18719	18720
NT2RP3003799	C-NT2RP3003799	18721	18722
NT2RP3003800	C-NT2RP3003800	18723	18724
NT2RP3003828	C-NT2RP3003828	18725	18726
NT2RP3003932	C-NT2RP3003932	18727	
NT2RP3003992	C-NT2RP3003992	18728	18729
NT2RP3004013	C-NT2RP3004013	18730	18731
NT2RP3004028	C-NT2RP3004028	18732	18733
NT2RP3004041	C-NT2RP3004041	18734	18735
NT2RP3004051	C-NT2RP3004051	18736	18737
NT2RP3004078	C-NT2RP3004078	18738	18739
NT2RP3004093	C-NT2RP3004093	18740	18741
NT2RP3004095	C-NT2RP3004095	18742	18743
NT2RP3004125	C-NT2RP3004125	18744	18745
NT2RP3004148	C-NT2RP3004148	18746	18747
NT2RP3004155	C-NT2RP3004155	18748	18749
NT2RP3004189	C-NT2RP3004189	18750	18751
NT2RP3004332	C-NT2RP3004332	18752	18753
NT2RP3004349	C-NT2RP3004349	18754	
NT2RP4000035	C-NT2RP4000035	18755	
NT2RP4000049	C-NT2RP4000049	18756	
NT2RP4000102	C-NT2RP4000102	18757	
NT2RP4000167	C-NT2RP4000167	18758	18759
NT2RP4000515	C-NT2RP4000515	18760	
NT2RP4000517	C-NT2RP4000517	18761	
NT2RP4000519	C-NT2RP4000519	18762	18763
NT2RP5003512	C-NT2RP5003512	18764	18765
OVARC1000092	C-OVARC1000092	18766	
OVARC1000533	C-OVARC1000533	18767	

(1016)

特開 2002-191363

2029

2030

OVARC1000678	C-OVARC1000678	18768	
OVARC1000689	C-OVARC1000689	18769	18770
OVARC1000802	C-OVARC1000802	18771	
OVARC1000890	C-OVARC1000890	18772	18773
OVARC1000945	C-OVARC1000945	18774	18775
OVARC1001072	C-OVARC1001072	18776	18777
OVARC1001117	C-OVARC1001117	18778	18779
OVARC1001200	C-OVARC1001200	18780	18781
OVARC1001244	C-OVARC1001244	18782	18783
OVARC1001329	C-OVARC1001329	18784	18785
OVARC1001341	C-OVARC1001341	18786	18787
OVARC1001376	C-OVARC1001376	18788	
OVARC1001496	C-OVARC1001496	18789	18790
OVARC1001873	C-OVARC1001873	18791	
PLACE1000007	C-PLACE1000007	18792	18793
PLACE1000547	C-PLACE1000547	18794	18795
PLACE1001036	C-PLACE1001036	18796	18797
PLACE1001076	C-PLACE1001076	18798	
PLACE1001118	C-PLACE1001118	18799	18800
PLACE1001366	C-PLACE1001366	18801	18802
PLACE1001608	C-PLACE1001608	18803	18804
PLACE1002004	C-PLACE1002004	18805	18806
PLACE1002256	C-PLACE1002256	18807	
PLACE1002437	C-PLACE1002437	18808	18809
PLACE1002591	C-PLACE1002591	18810	18811
PLACE1002665	C-PLACE1002665	18812	18813
PLACE1003864	C-PLACE1003864	18814	18815
PLACE1004793	C-PLACE1004793	18816	18817
PLACE1004913	C-PLACE1004913	18818	18819
PLACE1004979	C-PLACE1004979	18820	
PLACE1005052	C-PLACE1005052	18821	18822
PLACE1005055	C-PLACE1005055	18823	18824
PLACE1005128	C-PLACE1005128	18825	18826
PLACE1005162	C-PLACE1005162	18827	18828
PLACE1005176	C-PLACE1005176	18829	18830
PLACE1005467	C-PLACE1005467	18831	18832
PLACE1005584	C-PLACE1005584	18833	18834
PLACE1005611	C-PLACE1005611	18835	18836
PLACE1005802	C-PLACE1005802	18837	
PLACE1005850	C-PLACE1005850	18838	
PLACE1005898	C-PLACE1005898	18839	18840
PLACE1005932	C-PLACE1005932	18841	
PLACE1006129	C-PLACE1006129	18842	18843
PLACE1006360	C-PLACE1006360	18844	
PLACE1006795	C-PLACE1006795	18845	
PLACE1006878	C-PLACE1006878	18846	18847
PLACE1007557	C-PLACE1007557	18848	
PLACE1007807	C-PLACE1007807	18849	18850
PLACE1008181	C-PLACE1008181	18851	
PLACE1008426	C-PLACE1008426	18852	18853

(1017)

特開 2 0 0 2 - 1 9 1 3 6 3

2031

2032

PLACE1008941	C-PLACE1008941	18854	18855
PLACE1009935	C-PLACE1009935	18856	18857
PLACE1010310	C-PLACE1010310	18858	18859
PLACE1011891	C-PLACE1011891	18860	
PLACE1011896	C-PLACE1011896	18861	18862
PLACE2000003	C-PLACE2000003	18863	
PLACE2000132	C-PLACE2000132	18864	18865
PLACE2000170	C-PLACE2000170	18866	
PLACE2000335	C-PLACE2000335	18867	18868
PLACE3000124	C-PLACE3000124	18869	18870
PLACE3000158	C-PLACE3000158	18871	
PLACE3000207	C-PLACE3000207	18872	
PLACE3000221	C-PLACE3000221	18873	18874
PLACE3000271	C-PLACE3000271	18875	18876
PLACE3000304	C-PLACE3000304	18877	
PLACE3000322	C-PLACE3000322	18878	18879
PLACE3000341	C-PLACE3000341	18880	
PLACE3000373	C-PLACE3000373	18881	18882
PLACE3000399	C-PLACE3000399	18883	18884
PLACE3000401	C-PLACE3000401	18885	
PLACE3000402	C-PLACE3000402	18886	
PLACE3000406	C-PLACE3000406	18887	18888
PLACE3000475	C-PLACE3000475	18889	
PLACE4000063	C-PLACE4000063	18890	18891
PLACE4000093	C-PLACE4000093	18892	
PLACE4000100	C-PLACE4000100	18893	18894
PLACE4000247	C-PLACE4000247	18895	18896
PLACE4000250	C-PLACE4000250	18897	18898
PLACE4000252	C-PLACE4000252	18899	18900
PLACE4000259	C-PLACE4000259	18901	18902
PLACE4000320	C-PLACE4000320	18903	
PLACE4000344	C-PLACE4000344	18904	
PLACE4000367	C-PLACE4000367	18905	18906
PLACE4000401	C-PLACE4000401	18907	18908
PLACE4000411	C-PLACE4000411	18909	18910
PLACE4000487	C-PLACE4000487	18911	18912
PLACE4000494	C-PLACE4000494	18913	18914
PLACE4000521	C-PLACE4000521	18915	18916
PLACE4000548	C-PLACE4000548	18917	18918
SKNMC1000013	C-SKNMC1000013	18919	18920
SKNMC1000091	C-SKNMC1000091	18921	18922
THYRO1000343	C-THYRO1000343	18923	18924
THYRO1000569	C-THYRO1000569	18925	18926
THYRO1001142	C-THYRO1001142	18927	
THYRO1001189	C-THYRO1001189	18928	18929
THYRO1001320	C-THYRO1001320	18930	18931
THYRO1001537	C-THYRO1001537	18932	18933
THYRO1001721	C-THYRO1001721	18934	18935
THYRO1001828	C-THYRO1001828	18936	18937
Y79AA1000346	C-Y79AA1000346	18938	18939

(1018)

特開 2002-191363

2033

2034

Y79AA1001167	C-Y79AA1001167	18940	
Y79AA1001384	C-Y79AA1001384	18941	18942
Y79AA1001875	C-Y79AA1001875	18943	
Y79AA1002103	C-Y79AA1002103	18944	18945
HEMBA1000290	C-HEMBA1000290	18946	
HEMBA1001196	C-HEMBA1001196	18947	18948
HEMBA1006650	C-HEMBA1006650	18949	18950
HEMBA1006796	C-HEMBA1006796	18951	18952
HEMBB1000337	C-HEMBB1000337	18953	18954
HEMBB1001619	C-HEMBB1001619	18955	
MAMMA1000270	C-MAMMA1000270	18956	
MAMMA1000559	C-MAMMA1000559	18957	
MAMMA1000940	C-MAMMA1000940	18958	
MAMMA1002545	C-MAMMA1002545	18959	
MAMMA1002972	C-MAMMA1002972	18960	18961
NT2RP2001440	C-NT2RP2001440	18962	18963
NT2RP3002770	C-NT2RP3002770	18964	18965
NT2RP3003138	C-NT2RP3003138	18966	18967
NT2RP3004470	C-NT2RP3004470	18968	
OVARC1000891	C-OVARC1000891	18969	
PLACE1001545	C-PLACE1001545	18970	18971
PLACE1003383	C-PLACE1003383	18972	
PLACE1005549	C-PLACE1005549	18973	18974
PLACE1008455	C-PLACE1008455	18975	
PLACE4000131	C-PLACE4000131	18976	18977
PLACE4000261	C-PLACE4000261	18978	18979
THYR01001602	C-THYR01001602	18980	
HEMBA1006092	C-HEMBA1006092	18981	
HEMBA1006406	C-HEMBA1006406	18982	
HEMBB1000790	C-HEMBB1000790	18983	
HEMBB1000917	C-HEMBB1000917	18984	
HEMBB1002280	C-HEMBB1002280	18985	
MAMMA1000802	C-MAMMA1000802	18986	
MAMMA1001322	C-MAMMA1001322	18987	
MAMMA1002597	C-MAMMA1002597	18988	
MAMMA1002868	C-MAMMA1002868	18989	
NT2RP2003161	C-NT2RP2003161	18990	18991
NT2RP2003339	C-NT2RP2003339	18992	
NT2RP3001282	C-NT2RP3001282	18993	18994
PLACE1001761	C-PLACE1001761	18995	18996
PLACE1004491	C-PLACE1004491	18997	
PLACE1004686	C-PLACE1004686	18998	
PLACE1005574	C-PLACE1005574	18999	
PLACE1006382	C-PLACE1006382	19000	
PLACE1006792	C-PLACE1006792	19001	
PLACE3000455	C-PLACE3000455	19002	19003
PLACE4000230	C-PLACE4000230	19004	19005
THYR01000916	C-THYR01000916	19006	
HEMBA1000327	C-HEMBA1000327	19007	19008
HEMBB1000637	C-HEMBB1000637	19009	19010

2035		(1019)		2036	
HEMBB1001967	C-HEMBB1001967	19011			
MAMMA1000266	C-MAMMA1000266	19012			
NT2RP2002979	C-NT2RP2002979	19013			
PLACE1007866	C-PLACE1007866	19014	19015		
PLACE3000350	C-PLACE3000350	19016	19017		
PLACE4000156	C-PLACE4000156	19018	19019		
THYR01001637	C-THYR01001637	19020	19021		
MAMMA1002215	C-MAMMA1002215	19022	19023		
MAMMA1002721	C-MAMMA1002721	19024			
NT2RP2002070	C-NT2RP2002070	19025			

【0823】

【表352】滑膜細胞、TNF存在下で培養した滑膜細胞の各cDNAの発現（表中には実施例で説明の無いクローンも含まれている）

表中、Synoviocyteは滑膜細胞を、Synoviocyte_TNFはTNF処理した滑膜細胞を示す。また、exp1、exp2、およびex*

*p3は、それぞれn=3の測定結果を表している。ttest vs TNFは、滑膜細胞とTNF処理した滑膜細胞との有意差検定の結果を示す。TNF処理した滑膜細胞における発現量の増加を+で、減少を-で増減として示す。有意差検定の結果を、*P<0.05、および**P<0.01で示した。

Clone	Synoviocyte			Synoviocyte_TNF			t test	
	exp.1	exp.2	exp.3	exp.1	exp.2	exp.3	vs TNF	増減
GAPDH(Cr1)	0.4	0.8	0.89	0.9	1	1.15		
β actin(Cr2)	385.94	262.23	582.98	443.28	422.61	573.47		
ADRG11000005	2.72	2.97	4.46	7.27	7.45	3.51		
ADRG11000007	4.36	5.19	9.58	20.78	19.59	18.29	**	+
ADRG11000009	0.99	1.25	1.64	2.16	4.08	2.02		
ADRG11000011	1.98	3.56	5.24	22.22	23.49	19.81	**	+
ADRG11000027	0.79	1.22	1.66	2.82	4.99	1.9		
ADRG11000058	4.12	7.08	26.9	62.55	67.32	49.15	**	+
ADRG11000069	1.91	1.68	2.47	14.19	14.54	13.74	**	+
ADRG11000077	1.98	2	2.54	5.5	2.9	4.16		
ADRG11000092	2.99	4.79	12.53	21.46	22.09	26.19	**	+
ADRG11000099	2.77	4.79	12.85	23.61	24.02	25.56	**	+
ADRG11000136	20.49	27.18	31.85	62.44	40.69	48.29	*	+
ADRG11000147	2.09	2.58	5.47	5.69	7.52	3.85		
ADRG11000159	1.51	1.77	3.07	3.4	4.71	2.59		
ADRG11000160	2.42	4.34	6.89	8.08	7.24	7.06		
ADRG11000171	0.95	1.11	1.64	1.89	2.69	1.87		
ADRG11000181	0.64	1.37	1.74	3.99	4.27	3.89	**	+
BGG111000015	2.13	3.89	5.02	10.49	11.35	9.14	**	+
BGG111000016	27.77	35.71	52.17	57.18	48.51	63.57		
BGG111000017	1.29	3.19	3.14	3.24	3.65	2.34		
BGG111000022	4.72	4.45	6.75	10.71	5.56	8.27		
BGG111000031	4.47	6.58	8.77	14.79	11.63	10.04	*	+
BGG111000042	9.55	11.29	20.54	23.39	18.75	20.23		
BGG111000046	8.56	9.77	17.04	34.24	30.76	25.79	**	+
BNCH41000020	246.16	211.77	380.83	658.32	647.37	559.16	**	+
BNCH41000025	4.31	3.12	6.92	11.4	13.1	15.01	**	+
BNCH41000026	2.71	4.77	7.53	4.45	7.17	6.23		

(1020)

特開 2002-191363

2037

2038

BNGH41000027	11.52	13.5	12.69	20.62	12.48	24.91		
BNGH41000035	23.02	25.91	36.46	51.05	31.83	41.67		
BNGH41000037	2.7	5.21	6.72	12.95	8.98	8.59	*	+
BNGH41000042	14.55	16.06	22.84	49.62	37.57	36.25	**	+
BNGH41000048	3.92	6.27	25.68	66.19	74.4	66.21	**	+
BNGH41000056	0.74	1.75	3.26	5.28	7.34	3.75	*	+
BNGH41000087	3.36	4.08	5.19	5.59	8.15	3.01		
BNGH41000091	0.18	1.45	2.47	2.72	3.4	2.14		
BNGH41000157	6.93	7.99	6.23	13.37	10.28	9.98	*	+
BNGH41000169	1.09	1.53	2.99	2.77	4.23	2.59		
BNGH41000181	3.5	4.06	7.5	5.71	6.81	6.09		
BNGH41000198	1.32	2.3	4.35	2.06	2.55	2.22		
BNGH41000219	2.29	3.91	5.61	12.4	13.73	10.76	**	+
BNGH41000229	9.65	9.99	12.99	18.34	18.92	18.94	**	+
BNGH41000237	8.4	12.99	12.61	27.63	11.26	13.45		
BNGH41000238	1.56	2.59	6.77	3.45	4.55	3.32		
BNGH41000243	5.56	8.95	6.71	15.03	12.55	16.36	**	+
BNGH41000270	2.94	2.77	2.88	3.67	3.99	3.74	**	+
BRAWH1000004	1	2.19	6.99	6.45	8.36	6		
BRAWH1000018	1.8	2.24	5.06	4.43	6.95	5.24		
BRAWH1000021	1.33	2.73	4.81	4.16	5.85	5.21		
BRAWH1000027	0.58	1.7	1.62	2.39	3.65	2.63	*	+
BRAWH1000029	2.32	3.63	6.21	6.03	6.73	4.81		
BRAWH1000040	4.68	4.98	8.01	7.28	7.2	8.67		
BRAWH1000050	11.04	10.47	43.79	51.7	73.7	60.92	*	+
BRAWH1000051	2.14	0.63	2.71	2.25	4.43	1.04		
BRAWH1000060	7.84	8.07	48.26	59.16	66.12	63.86	*	+
BRAWH1000075	1.85	1.86	2.98	2.07	4.4	2.34		
BRAWH1000081	1.88	2.78	7.19	5.9	10.82	7.4		
BRAWH1000084	30.23	30.57	65.21	235.81	180.86	211.35	**	+
BRAWH1000095	1.38	2.47	4.51	3	4.78	2.67		
BRAWH1000096	1.37	2.89	4.71	3.7	4.8	5.17		
BRAWH1000097	3.32	3.27	10.74	9.24	10.62	7.75		
BRAWH1000100	4.77	5.19	7.69	6.98	7.06	7.28		
BRAWH1000101	12	12.04	36.52	46.19	41.09	50.21	*	+
BRAWH1000104	1.37	0.92	4.33	1.47	4.47	2.41		
BRAWH1000107	0.62	1.88	2.48	2.43	5.03	3.15		
BRAWH1000110	4.4	4.06	16.81	13.87	11.1	15.74		
BRAWH1000111	3.98	6.14	6.05	8.85	8.95	10.64	*	+
BRAWH1000135	4.95	4.91	7.7	7.37	9.42	9.98		
BRAWH1000190	2.22	3.84	5.07	4.66	7.16	4.99		
HEMBA1000005	5.91	6.44	11.97	17.55	22.88	18.65	*	+
HEMBA1000006	2.61	3.17	4.64	3.08	8.49	4.75		
HEMBA1000012	10.97	11.75	51.07	71.4	106.82	74.8	*	+
HEMBA1000020	50.65	49.12	113.3	197.41	293.79	216.89	*	+
HEMBA1000030	1.93	3.08	4.67	5.72	3.62	6.43		
HEMBA1000034	3.27	3.21	5.35	4.62	10.29	6.85		
HEMBA1000042	1.64	3.17	6	4.72	6.92	8.12		
HEMBA1000045	7.13	9.44	11.07	9.55	14.43	10.44		
HEMBA1000046	1.14	2.24	2.77	3.73	5.3	4.34	*	+

(1021)

特開2002-191363

2039

2040

HEMBA1000047	1.17	1.99	3.83	2.98	4.47	3.78		
HEMBA1000048	3.76	4.75	3.64	8.73	12.38	5.48		
HEMBA1000050	0.48	1.77	1.78	1.4	3.39	1.46		
HEMBA1000053	1.64	1.28	2.57	4.68	4.03	3.99	**	+
HEMBA1000060	1.88	2.71	4.51	7.29	7.94	9.74	**	+
HEMBA1000072	52.79	53.46	135.73	165.97	221.75	230.97	*	+
HEMBA1000073	16.54	11.43	27.32	22.4	36.09	33.78		
HEMBA1000076	5.06	5.33	9.77	12.16	10.46	11.15	*	+
HEMBA1000084	4.75	4.46	20.71	30.15	43.67	33.92	*	+
HEMBA1000087	0.51	1	3.32	0.65	2.82	1.61		
HEMBA1000088	1.98	2.97	4.6	6.2	9.87	8.46	*	+
HEMBA1000091	6.36	5.4	17.56	30.15	44.04	35.43	**	+
HEMBA1000111	1.52	1.77	3.63	5.29	6.65	6.4	**	+
HEMBA1000121	0.86	1.17	3.58	3.52	4.47	5.83		
HEMBA1000128	1.52	2.99	6.04	4.28	6.05	5.93		
HEMBA1000129	2.04	1.81	3.95	2.66	3.26	3.32		
HEMBA1000141	2.31	3.45	5.98	3.56	6.67	5.6		
HEMBA1000146	0.84	1.29	2.96	1.93	4.98	3.52		
HEMBA1000150	3.34	3.29	10.65	8.27	11.97	9.46		
HEMBA1000154	25.17	29.21	82.33	128.3	134.42	139.59	**	+
HEMBA1000156	3.28	4	5.87	8.69	6.19	6.13		
HEMBA1000158	7.98	10.04	12.52	16.99	15.47	12.88		
HEMBA1000168	1.21	2.2	4.11	5.7	7.3	5.21	*	+
HEMBA1000180	0.4	2.04	2.87	2.86	4.06	2.05		
HEMBA1000185	1.65	3.84	4.88	7.5	9.46	9.07	**	+
HEMBA1000188	1.37	1.64	3.31	4.94	4.19	3.35		
HEMBA1000193	1.53	0.66	3.16	2.68	4.33	2.5		
HEMBA1000194	2.18	2.95	5.68	9.11	8.74	8.83	**	+
HEMBA1000201	2.6	4.47	9.74	13.45	14.8	14.65	*	+
HEMBA1000213	1.33	1.95	2.76	2.08	4.49	3.7		
HEMBA1000216	1.26	1.82	3.27	2.92	5.2	3.47		
HEMBA1000227	0.99	2.27	2.38	3.28	4.21	1.83		
HEMBA1000231	1.5	1.97	5	7.56	7.19	6.16	*	+
HEMBA1000237	4.5	6.13	9.14	14.79	18.3	14.71	**	+
HEMBA1000243	0.6	1.89	3.01	4.3	4.34	3.67	*	+
HEMBA1000244	1.54	2.45	3.78	6.08	5.58	3.36		
HEMBA1000251	1.15	1.92	2.97	2.26	4.59	3.22		
HEMBA1000254	0.69	1.8	4.81	3.57	4.97	3.58		
HEMBA1000264	0.84	2.28	3.01	2.84	3.23	3.12		
HEMBA1000269	1.9	2.34	3.69	4.41	4.09	2.51		
HEMBA1000275	5.31	4.29	8.03	7.96	12.04	8.54		
HEMBA1000280	1.43	0.83	2.19	3.3	4.08	4	**	+
HEMBA1000282	1.15	1.01	4.23	6.29	7.01	5.46	*	+
HEMBA1000287	2.86	3.19	4.45	5.81	6.04	6.37	**	+
HEMBA1000288	1.37	2.23	6.13	3.51	6.02	3.85		
HEMBA1000290	1.01	2.17	4.11	2.46	3.26	2.73		
HEMBA1000296	2.4	3.66	5.49	6.15	6.55	5.84		
HEMBA1000300	1.22	2.73	6.6	7.64	8.88	7.23		
HEMBA1000302	0.93	2.17	2.86	3.04	3.74	1.97		
HEMBA1000303	1.36	2.15	3.57	4.13	4.43	3		

(1022)

特開 2002-191363

2041

2042

HEMBA1000304	1.06	1.99	4.26	5.51	7.28	4.87	*	+
HEMBA1000307	1.21	1.73	2.65	4.4	5.64	2.99	*	+
HEMBA1000312	6	8.7	10.77	13.2	9.18	9.65		
HEMBA1000318	1.5	4.22	3.25	5.39	6.05	4.49		
HEMBA1000327	2.18	3.7	3.34	10.58	6.06	6.02	*	+
HEMBA1000333	0.68	2.75	4.33	3.12	4.74	2.98		
HEMBA1000338	1.61	2.84	5.33	5.8	5.78	4.32		
HEMBA1000343	1.79	3.5	3.69	5.55	6.7	3.99		
HEMBA1000349	0.97	1.52	3.24	3.9	5.37	4.09	*	+
HEMBA1000351	1.6	2.06	5.75	4.8	6.22	5.24		
HEMBA1000355	1.52	3.09	4.09	3.78	5.14	3.59		
HEMBA1000356	9.3	10.42	14.39	26.93	22.26	24.97	**	+
HEMBA1000357	1.88	2.11	4.76	3.81	5.7	4.62		
HEMBA1000366	1.67	1.94	3.83	3.14	4.75	3.28		
HEMBA1000369	1.87	2.94	5.17	2.82	5.2	4.56		
HEMBA1000370	2.45	3.4	4.63	3.75	5.34	3.6		
HEMBA1000376	3.64	4.55	14.48	26.69	29.98	28.36	**	+
HEMBA1000387	2.95	3.19	6.2	7.85	7.62	8.15	*	+
HEMBA1000389	2.88	3.74	8.83	14.4	10.9	13.61	*	+
HEMBA1000390	1.86	2.27	3.5	4.28	4.98	3.95	*	+
HEMBA1000392	1.49	1.4	3.06	2.58	3.78	1.94		
HEMBA1000396	1.82	2.16	3.45	3.43	4.93	3.34		
HEMBA1000411	1.01	1.41	4.49	1.94	4.41	2.21		
HEMBA1000418	2.85	3.21	4.41	7.75	6.81	5.17	*	+
HEMBA1000422	0.99	1.89	2.14	2.64	4.03	2.89		
HEMBA1000428	0.36	2.43	3.09	2.58	3.31	2.75		
HEMBA1000434	0.54	2.19	2.93	2.11	3.6	2.69		
HEMBA1000442	0.82	2.2	3.37	2.13	3.8	2.28		
HEMBA1000443	1.19	1.9	3.12	2.99	6.28	3.59		
HEMBA1000446	38.48	43.56	75.05	56.34	60.86	69.87		
HEMBA1000456	5.19	4.41	6.5	7.45	5.62	8.77		
HEMBA1000459	1.95	2.11	4.24	3.46	6.17	5.55		
HEMBA1000460	7.46	7.84	8.87	13.59	12.54	18.45	*	+
HEMBA1000462	2.11	3.51	5.04	6.05	5.16	7.49		
HEMBA1000464	1.33	0.96	1.73	1.69	2.74	2.53		
HEMBA1000468	1.25	1.44	2.43	1.69	3.48	2.22		
HEMBA1000469	2.89	3.37	8.1	5.42	8.81	8.01		
HEMBA1000477	2.87	3.03	7.4	5.41	9.68	6.83		
HEMBA1000481	29.67	31.97	31.95	42.76	52.75	25.82		
HEMBA1000488	1.75	2.43	2.96	3.11	5.9	3		
HEMBA1000490	1.34	2	3.49	4.41	3.7	2.88		
HEMBA1000491	1.21	1.71	2.85	4.24	4.99	5.97	*	+
HEMBA1000498	2.12	3.21	4.55	4.39	7.76	5.94		
HEMBA1000501	2.22	3.36	6.25	6.44	8.93	9.74	*	+
HEMBA1000504	2.93	3.18	4.82	3.63	5.37	3.83		
HEMBA1000505	0.81	1.97	3.33	2.72	5.1	3.58		
HEMBA1000507	1.02	2.24	5.29	4.17	8.62	7		
HEMBA1000508	2.25	2.3	7.65	4.84	8.57	6.64		
HEMBA1000518	1.38	0.96	0.98	1.89	2.97	1.8	*	+
HEMBA1000519	9.5	7.28	15.97	19.28	20.99	19.72	*	+

(1023)

特開 2002-191363

2043

2044

HEMBA1000520	0.45	1.12	1.18	1.94	4.83	4.3	*	+
HEMBA1000523	2.32	1.88	3.22	3.48	5.33	3.65		
HEMBA1000531	1.39	1.46	2.44	2.67	5.34	4.63	*	+
HEMBA1000534	0.55	0.95	2.97	6.63	11.62	10.39	**	+
HEMBA1000538	0.51	1.08	2.31	12.58	21.02	13.18	**	+
HEMBA1000540	2.8	3.11	6.06	5.82	10.38	6.39		
HEMBA1000542	9.16	7.79	43.94	62.25	95.7	81.15	*	+
HEMBA1000545	1.51	2.31	1.65	3.19	4.29	3.7	**	+
HEMBA1000547	2.99	3.12	4.94	4.94	5.3	4.97		
HEMBA1000551	2.32	1.99	9.54	4.68	7.33	9.81		
HEMBA1000555	3.81	3.23	6.39	5.03	6.43	8.08		
HEMBA1000557	2.16	2.06	6.07	3.98	6.46	5.06		
HEMBA1000561	1.71	2.9	4.9	1.63	4.39	3.67		
HEMBA1000563	1.73	1.85	4.09	2.72	3.94	2.83		
HEMBA1000567	1.02	1.01	1.67	1.21	2.59	1.92		
HEMBA1000568	2.19	2.5	6.09	7.62	6.65	6.84		
HEMBA1000569	1.3	2.8	3.02	2.18	6.47	2.3		
HEMBA1000575	3.73	4.91	10.84	10.19	15.17	13.08		
HEMBA1000588	1.75	2.49	4.16	3.12	5.5	3.83		
HEMBA1000590	0.59	1.02	2.06	2.24	2.53	1.35		
HEMBA1000591	3.17	3.3	5.18	10.84	12.16	9.8	**	+
HEMBA1000592	4.2	5.19	7.77	13.85	14.94	11.78	**	+
HEMBA1000594	1.95	1.97	3.16	4	5.86	4.94	*	+
HEMBA1000604	1.19	3.37	3.48	5.41	10.91	5.29		
HEMBA1000607	2.83	5.09	12.7	15.52	18.13	20.66	*	+
HEMBA1000608	0.9	2.34	2.46	2.6	5.5	2.31		
HEMBA1000622	0.96	2.19	3.55	3.61	5.24	3.8		
HEMBA1000634	17.56	22.96	30.36	71.62	60.59	51.59	**	+
HEMBA1000636	4.59	3.95	6.78	15.48	12.35	12.73	**	+
HEMBA1000637	0.93	0.48	2.58	2.42	3.19	2.21		
HEMBA1000655	1.33	2.11	4.84	6.91	5.57	6.31	*	+
HEMBA1000657	1.35	1.78	3.24	4.89	5.28	3.26	*	+
HEMBA1000662	1.3	2.42	2.73	2.52	3.78	2.72		
HEMBA1000664	0.94	1.6	2.87	3.11	4.63	2.94		
HEMBA1000671	2.96	3.84	11.68	21.25	18.69	15.76	*	+
HEMBA1000673	1.46	2.23	4.76	7.44	7.49	5.51	*	+
HEMBA1000675	4.18	3.09	4.54	8.18	7.19	8.04	**	+
HEMBA1000678	2.23	2.7	4.47	5.03	7.16	5.16		
HEMBA1000682	3.4	4.64	8.41	13.76	13.69	14.29	**	+
HEMBA1000686	2.73	3.88	4.83	6.23	6.6	5.32	*	+
HEMBA1000702	1.56	2.07	5.25	4.15	5.78	4.32		
HEMBA1000705	0.65	1.71	3.43	2.34	3.21	1.64		
HEMBA1000713	3.31	5.6	6.12	6.94	5.86	5.47		
HEMBA1000718	2.14	2.7	5.25	6.11	5.09	5.95		
HEMBA1000719	9.64	12.27	17.77	16.64	15.52	15.64		
HEMBA1000722	1.97	1.7	3.6	6.55	6.45	5.02	**	+
HEMBA1000726	2.2	2.23	5.12	9.4	8.77	9.36	**	+
HEMBA1000727	4.09	5.35	6.41	5.13	9.08	8.37		
HEMBA1000732	1.22	2.74	4.21	4.93	5.58	4.42		
HEMBA1000736	1.56	2.15	3.24	4.11	5.19	4.62	*	+

(1024)

特開 2002-191363

2045

2046

HEMBA1000743	1.25	2.72	3.41	5.05	4.88	4.16	*	+
HEMBA1000745	1.59	2.47	3.64	4.88	5.33	3.49		
HEMBA1000747	1.19	1.59	2.56	2.35	3.12	1.49		
HEMBA1000748	1.67	1.51	4.85	5.11	6.08	4.81		
HEMBA1000749	1.14	2.04	5.69	5.98	5.91	5.96		
HEMBA1000752	1.4	2.3	4.38	3.69	4.53	3.85		
HEMBA1000753	2.56	4.21	6.53	7.98	8.59	4.93		
HEMBA1000757	1.95	2.95	3.27	6.33	6.68	5.94	**	+
HEMBA1000760	3.71	3.81	6.62	6.96	7.03	6.89		
HEMBA1000769	1.99	2.36	5.17	3.48	5.87	2.85		
HEMBA1000773	1	2.32	3.07	2.17	3.18	1.4		
HEMBA1000774	2.69	2.76	6.37	6.29	7.77	5.22		
HEMBA1000780	1.12	2.33	3.66	2.7	4.78	3.29		
HEMBA1000783	1.32	2.39	4.1	2.78	7.73	2.57		
HEMBA1000791	2.07	2.4	6.39	4.97	10.17	7.84		
HEMBA1000793	12.73	12.73	17.88	19.93	17.49	16.69		
HEMBA1000802	1.57	1.65	2.59	2.07	4.41	1.1		
HEMBA1000813	38.24	35.83	34.83	54.63	42.38	53.94	*	+
HEMBA1000817	2.63	3.82	5.44	5.12	7.02	5.49		
HEMBA1000822	1.83	2.89	4.1	4.42	5.76	3.91		
HEMBA1000827	2.26	2.74	6.45	9.31	7.75	6.94	*	+
HEMBA1000833	3.1	4.46	7.31	8.06	4.49	4.85		
HEMBA1000835	12.53	15.55	75.61	94.51	110.02	86.95	*	+
HEMBA1000843	1.21	2.2	4.6	3.32	5.63	4.93		
HEMBA1000851	2.13	1.26	3.5	2.7	5.61	2.74		
HEMBA1000852	1.95	1.83	5.5	3.52	5.49	3.83		
HEMBA1000867	0.85	2.79	4.72	2.77	5.39	3.07		
HEMBA1000869	0.58	1.29	2.51	2.84	3.97	2.38		
HEMBA1000870	2.56	2.97	2.59	3.39	5.16	5.49	*	+
HEMBA1000872	1.44	2.87	4.01	4.31	4.14	4.34		
HEMBA1000875	1.89	3.09	5	3.8	4.38	3.77		
HEMBA1000876	1.75	3.36	4.64	3.9	6.21	4.9		
HEMBA1000907	1.99	2.47	3.81	3.21	7.15	5.53		
HEMBA1000908	0.81	2.06	3.85	2	5.43	1.98		
HEMBA1000910	1.97	1.61	3.71	3.35	5.25	2.98		
HEMBA1000918	0.76	1.34	4.37	4.93	6.54	6.95	*	+
HEMBA1000919	0.86	1.97	2.19	2.49	3.07	3.07		
HEMBA1000934	2.5	2.56	1.16	2.14	3.51	2.5		
HEMBA1000935	1.46	1.62	4.21	2.08	5.15	3.64		
HEMBA1000940	1.98	3.08	3.1	2.52	9.96	5.72		
HEMBA1000942	2.31	2.27	4.77	4.81	7.75	6.69		
HEMBA1000943	0.58	1.25	2.28	1.83	3.38	2.18		
HEMBA1000946	3.63	4.04	4.54	6.87	14.9	8.4		
HEMBA1000960	2.63	3.48	9.97	10.24	12.79	10.7		
HEMBA1000962	1.99	2.18	2.01	4.43	3.83	4.56	**	+
HEMBA1000968	1.73	1.86	4.7	4.1	4.83	4.66		
HEMBA1000971	1.75	2.51	2.9	4.18	5.27	5.71	**	+
HEMBA1000972	1.45	1.57	3.83	2.63	4.44	3.49		
HEMBA1000974	1.69	2.69	6.33	7.39	9.35	8.82	*	+
HEMBA1000975	0.9	1.83	4.17	3.31	5.54	5.12		

(1025)

特開 2002-191363

2047

2048

HEMBA1000979	1.45	1.69	3.98	2.55	6.12	3.93		
HEMBA1000981	4.21	6.9	9.5	11.75	13.27	14.72	*	+
HEMBA1000983	1.94	1.45	3.01	3.89	4.53	4.15	*	+
HEMBA1000985	1.58	0.92	2.75	1.73	3.28	2.79		
HEMBA1000986	1.2	1.48	2.47	3.61	4.91	4.26	**	+
HEMBA1000991	1.56	1.86	3.8	3.11	5.05	5.96		
HEMBA1001007	0.89	1.08	4.08	1.84	3.89	2.71		
HEMBA1001008	3.64	3.41	5.86	3.89	7.89	4.95		
HEMBA1001009	0.89	1.3	3.07	1.58	3.83	1.81		
HEMBA1001014	3.54	4.39	9.91	11.82	15.38	14.12	*	+
HEMBA1001017	4.21	2.82	5.6	6.04	5.41	8.55		
HEMBA1001019	1.92	2.81	3.97	8.71	7.74	8.29	**	+
HEMBA1001020	1.23	2.71	2.3	2.84	5.05	3.6		
HEMBA1001021	1.07	1.62	2.89	3.13	5.24	2.63		
HEMBA1001022	2.29	2.25	4.35	6.33	8.57	3.81		
HEMBA1001024	0.31	1.14	2.16	2.87	3.97	1.26		
HEMBA1001026	0.42	1.52	1.86	2	3.22	2		
HEMBA1001043	1.43	2.46	2.38	4.63	5.28	4.25	**	+
HEMBA1001051	3.36	2.79	11.52	13.26	18.17	18.47	*	+
HEMBA1001052	0.86	2.15	2.18	1.75	3.58	2.48		
HEMBA1001059	5.62	9.28	26.25	40.62	56.12	43.49	*	+
HEMBA1001060	2.66	3.67	6.45	10.78	8.35	9.62	*	+
HEMBA1001064	2.12	2.87	3.3	6.04	6.48	4.69	**	+
HEMBA1001071	29.39	41.54	55.57	143.9	102.43	121.71	**	+
HEMBA1001077	2.37	1.77	5.21	5.36	6.66	3.96		
HEMBA1001078	2.18	2.6	5.91	13.3	13.21	11.09	**	+
HEMBA1001080	4.03	3.46	11.86	24.15	26.66	26.65	**	+
HEMBA1001084	1.27	2.37	2.9	5.07	5.88	5.13	**	+
HEMBA1001085	1.24	2.87	4.04	4.34	5.41	4.56		
HEMBA1001088	6.62	6	8.04	3.79	4.34	5.81		
HEMBA1001093	0.61	1.76	2.72	3.09	3.02	2.99		
HEMBA1001094	0.64	0.78	2.07	2.08	2.99	1.99		
HEMBA1001099	1.01	1.72	3	2.5	2.95	2.26		
HEMBA1001104	1.2	1.75	2.63	3.64	8.04	3.3		
HEMBA1001109	4.87	3.77	8.57	11.32	14.48	11.73	*	+
HEMBA1001114	44.68	41.2	93.35	141.87	145.19	167.76	**	+
HEMBA1001121	2.14	2.03	3.87	2.41	6	3.25		
HEMBA1001122	9.79	10	14.12	7.73	11.5	22.69		
HEMBA1001123	2.79	3.28	5.2	5.81	6.02	4.95		
HEMBA1001133	0.97	1.69	2.54	2.78	3.84	1.21		
HEMBA1001137	0.82	1.73	3.65	3.74	3.36	2.54		
HEMBA1001140	1.23	2.75	2.98	3.62	5.18	4.34	*	+
HEMBA1001144	4.12	3.41	9.06	14.13	14.12	13.96	**	+
HEMBA1001145	47.87	43.87	65.7	98.4	75.15	81.3	*	+
HEMBA1001158	7.55	9.5	11.62	13.02	7.58	12.5		
HEMBA1001172	1.44	2.85	4.37	5.32	5.77	5.17	*	+
HEMBA1001174	0.95	2.06	2.83	3.88	6.31	3.25		
HEMBA1001175	6.93	8.56	10.73	14.17	14.5	10.18		
HEMBA1001182	16.93	19.89	82.44	135.93	145.36	122.22	*	+
HEMBA1001184	1.41	1.24	2.45	1.85	3.03	1.47		

(1026)

特開 2002-191363

2049

2050

HEMBA1001192	1.72	1.75	4.01	5.65	5.17	3.98		
HEMBA1001196	2.31	3.63	7.61	9.43	10.51	8.97	*	+
HEMBA1001197	31.18	35.89	86.14	95.35	83.09	93.59		
HEMBA1001208	1.83	2.59	3	2.67	5.3	2.61		
HEMBA1001213	12.99	16.12	69.9	102.88	119.96	113.72	*	+
HEMBA1001214	1.39	3.11	4.36	5.14	7.04	4.62		
HEMBA1001221	1.63	1.62	3.66	2.06	4.19	1.89		
HEMBA1001225	1.06	2.66	3.53	1.44	3.43	1.52		
HEMBA1001226	4.76	4.65	11.94	13.58	15.58	14.92	*	+
HEMBA1001228	72.4	75.3	102.4	38.23	64.63	78.89		
HEMBA1001229	18	21.39	82.05	115.91	145.39	128.91	*	+
HEMBA1001235	3.58	4.11	6.48	7.31	6.7	10.2		
HEMBA1001238	2.46	2.49	7.23	4.6	6.94	4.74		
HEMBA1001242	15.36	14.03	91.45	92.81	94.02	90.34		
HEMBA1001247	4.41	4.36	12.46	12.48	14.07	15.62		
HEMBA1001253	8.79	11.4	61.56	77.17	102.24	94.81	*	+
HEMBA1001257	1.98	2.71	3.78	3.52	4.29	3.17		
HEMBA1001261	3.01	3.18	4.56	4.54	3.75	5.59		
HEMBA1001262	1.48	3.79	2.81	2.42	4.34	4.59		
HEMBA1001265	2.76	3.21	6.85	5.5	7.32	5.1		
HEMBA1001266	3.97	3.17	6.31	7.8	10.5	8.38	*	+
HEMBA1001269	15.98	10.36	12.79	22.69	24.71	25.21	**	+
HEMBA1001272	1.31	2.04	4.3	1.62	5.12	2.07		
HEMBA1001279	2.54	3.52	13.6	18.68	23.45	18.99	*	+
HEMBA1001281	16.58	20.99	40.84	47.71	59.04	45.72	*	+
HEMBA1001286	3.25	4.71	10.71	11.24	10.65	12.38		
HEMBA1001289	0.41	1.57	1.64	1.3	3.57	2.41		
HEMBA1001291	3.52	4.58	9.53	10.91	18.3	18.8	*	+
HEMBA1001294	2.01	1.81	4.6	4.04	7.73	5.12		
HEMBA1001296	3.4	3.52	4.37	3.77	5.94	5.22		
HEMBA1001297	2.88	3.61	5.51	4.81	6.88	5.38		
HEMBA1001299	2.49	2.9	6.21	6.45	8.84	7.74	*	+
HEMBA1001302	9.42	11.94	15.5	23.25	35.12	25.74	*	+
HEMBA1001303	1.8	1.99	2.61	3.57	3.8	3.3	**	+
HEMBA1001306	1.4	1.15	2.85	5.01	4.46	4.82	**	+
HEMBA1001308	3.43	4.37	16.7	16.31	18.28	21.75		
HEMBA1001310	1.93	1.71	4.17	2.38	6.26	3.28		
HEMBA1001312	10.09	10.35	17.42	20.51	24.71	21.67	*	+
HEMBA1001319	1.23	1.41	3.85	2.23	4.27	4.01		
HEMBA1001322	1.81	2.29	4.17	2.83	4.74	3.78		
HEMBA1001323	4.04	3.65	8.44	14.68	23.44	18.68	*	+
HEMBA1001326	8.79	7.35	10.15	12.24	13.62	15.04	*	+
HEMBA1001327	0.94	1.65	3.18	3.55	5.18	4.56	*	+
HEMBA1001330	1.59	2.22	6.96	7.36	9.28	9.64	*	+
HEMBA1001348	1.68	3.99	3.89	6.33	9.84	7.47	*	+
HEMBA1001350	5.28	4.16	6.34	7.24	13.17	10.12		
HEMBA1001351	15.37	14.99	17.64	37.37	49.52	25.96	*	+
HEMBA1001352	3.25	3.62	5.97	8.16	13.65	5.75		
HEMBA1001353	30.24	37.73	49.4	76.74	96.09	96.34	**	+
HEMBA1001358	13.98	9.73	17.96	30.89	27.69	30.6	**	+

(1027)

特開2002-191363

2051

2052

HEMBA1001361	1.7	3.24	4.96	4.18	6.08	6.06		
HEMBA1001364	0.8	1.71	2.4	1.47	4.11	2.95		
HEMBA1001375	3.45	2.77	5.75	5.71	5.83	6.32		
HEMBA1001377	2.81	3.16	7.36	5.37	7.98	7.89		
HEMBA1001383	0.25	1.64	2.61	1.26	2.47	1.84		
HEMBA1001387	1.81	2.15	3.66	1.94	5.14	2.47		
HEMBA1001388	1.52	1.78	5.07	2.01	4.61	3.49		
HEMBA1001390	34.61	34.52	66.57	67.03	50	56.4		
HEMBA1001391	1.65	2.77	4.83	4.32	7.98	3.82		
HEMBA1001398	1.98	2.87	7.47	7.24	10.42	8.29		
HEMBA1001405	1.17	2	3.87	2.99	5.3	2.61		
HEMBA1001406	2.01	3.27	3.75	5.35	6.62	4.33	*	+
HEMBA1001407	1.13	1.78	3.73	6.39	6.64	4.44	*	+
HEMBA1001411	1.44	2.81	4.47	7.2	6.35	6.04	*	+
HEMBA1001413	1.84	1.53	3.31	3.61	3.75	3.76		
HEMBA1001414	1.47	2.34	5.3	5.33	7.22	5.47		
HEMBA1001415	1.91	2.36	4.97	4.4	5.86	4.29		
HEMBA1001416	4.73	4.85	9.54	8.87	11.06	9.4		
HEMBA1001432	1.23	1.27	4.43	4.27	6.64	3.59		
HEMBA1001433	1.96	2.93	4.55	4.33	8.66	3.64		
HEMBA1001435	2.17	2.27	6.39	7.02	10.35	5.88		
HEMBA1001442	0.99	0.68	2.02	2.36	3.12	1.81		
HEMBA1001446	1.87	1.84	5.82	9.71	8.93	11.01	**	+
HEMBA1001450	2.35	2.32	11.22	8.61	10.08	6.34		
HEMBA1001454	3.08	4.25	9.69	13.64	10.73	11.82	*	+
HEMBA1001455	2.28	2.7	3.11	2.69	5.54	2.8		
HEMBA1001459	2.74	3.37	6.03	5.07	7.38	5.52		
HEMBA1001461	3.34	4.47	6.96	6.8	9.85	7.47		
HEMBA1001462	1.07	1.47	2.79	2.67	4.5	2.54		
HEMBA1001463	1.38	1.61	5.25	4.95	5.46	5.51		
HEMBA1001469	3.9	4.51	7.32	10.63	9.83	7.76	*	+
HEMBA1001473	4.56	3.49	8.25	7.52	10.34	6.53		
HEMBA1001477	2.14	1.59	4.64	3.41	5.75	2.59		
HEMBA1001478	2.46	2.8	3.77	2.95	3.73	2.55		
HEMBA1001480	4.15	6.8	8.96	11.64	11.87	8.48		
HEMBA1001483	1.9	1.64	5.71	6.6	8.22	7.23	*	+
HEMBA1001490	1.45	2.09	3.76	5.16	4.52	4.65	*	+
HEMBA1001495	56.8	53.41	123.27	193.11	133.65	132.04		
HEMBA1001497	2.06	1.98	7.47	4.81	8.45	5.85		
HEMBA1001510	3.99	4.23	15.22	11.46	13.7	14.56		
HEMBA1001515	1.45	2.33	4.02	3.73	6.11	3.04		
HEMBA1001517	1.6	2.21	4.6	5.26	5.4	4.6		
HEMBA1001522	1.56	2.72	3.77	3.61	6.37	2.43		
HEMBA1001526	2.19	2.97	4.97	4.05	4.38	3.59		
HEMBA1001533	3.19	2.86	6.23	6.83	7.76	4.64		
HEMBA1001547	7.26	5.37	13.69	5	7.96	6.19		
HEMBA1001552	7.12	4.72	17.79	16.12	16.3	16.05		
HEMBA1001553	41.67	45.48	66.23	57.2	47.01	79.81		
HEMBA1001557	2.24	2.93	5.15	5.81	8.33	4.59		
HEMBA1001563	1.69	2.4	4.56	3.66	6.99	4.76		

(1028)

特開2002-191363

2053

2054

HEMBA1001566	1.42	3.27	8.29	5.94	9.04	5.84		
HEMBA1001569	11.15	11.91	26.6	30.2	31.14	32.61	*	+
HEMBA1001570	3.25	4.61	10.2	9.19	10.25	9.53		
HEMBA1001579	3.63	4.4	7.77	9.26	7.4	9.93		
HEMBA1001581	2.79	3.33	10.95	8.81	12.09	9.08		
HEMBA1001582	3.22	3.18	6.68	6.35	6.84	4.03		
HEMBA1001585	2.7	3.07	4.52	4.4	5.6	3.5		
HEMBA1001589	1.82	2.31	3.63	4.39	6.19	3.78		
HEMBA1001595	13.06	15.57	19.7	13.25	13.29	14.02		
HEMBA1001604	1.96	2.67	3.64	3.76	6.53	2.82		
HEMBA1001608	5.58	7.09	16.17	14.14	16.46	14.43		
HEMBA1001615	113.28	90.33	205.41	240.97	118.65	165.59		
HEMBA1001620	3.71	5.56	10.54	12.22	12.24	11.46		
HEMBA1001621	0.76	2.13	3.42	1.76	3.44	2.97		
HEMBA1001635	2.32	2.13	3.41	3.55	4.9	2.85		
HEMBA1001636	1.9	1.93	4.01	3.34	5.33	2.97		
HEMBA1001640	3.07	3.31	13.65	10.96	15.01	10.74		
HEMBA1001647	8.92	8.44	57.38	88.92	112.42	87.46	*	+
HEMBA1001651	2.53	3.54	7.85	6.62	9.07	8.73		
HEMBA1001655	2.09	2.66	4.78	3.35	6.75	4.09		
HEMBA1001658	4.33	4.5	9.27	7.26	11.15	8.6		
HEMBA1001661	0.75	1.78	2.8	1.98	3.22	1.77		
HEMBA1001665	1.52	1.85	3.47	2.63	6.63	1.73		
HEMBA1001670	5.32	6.54	8.82	12.45	15.21	12.42	**	+
HEMBA1001672	2.49	3.06	5.9	4.28	7.62	3.39		
HEMBA1001673	8.23	10.76	13.22	20.04	19.39	15.65	*	+
HEMBA1001675	2.4	2.01	2.53	3.21	5.79	3.36		
HEMBA1001676	54.19	46.09	107.65	245.72	212.81	275.65	**	+
HEMBA1001678	9.46	10.2	21.87	23.65	19.51	27.88		
HEMBA1001680	4.58	4.89	12.32	9.39	10.95	11.65		
HEMBA1001681	1.71	2.44	5.75	6.25	9.11	6.36		
HEMBA1001684	1.89	2.74	6.26	4.32	7.57	6.98		
HEMBA1001695	1.48	2.08	3.42	2.3	4.76	3.15		
HEMBA1001702	1.54	2.96	3.55	2.36	7.57	3.09		
HEMBA1001709	1.23	1.8	3.51	3.21	4.87	3.5		
HEMBA1001711	1.29	1.98	2.83	2.99	2.45	3.18		
HEMBA1001712	0.92	1.55	2.56	2.13	3.02	2.24		
HEMBA1001714	10.37	10.82	19.06	23.54	22	23.8	*	+
HEMBA1001717	79.4	71.16	124.25	152.62	195.81	173.65	*	+
HEMBA1001718	1.95	2.12	7.32	5.99	6.59	5.26		
HEMBA1001723	3.43	3	10.19	9.09	12.53	9.64		
HEMBA1001731	1.3	1.36	3.27	1.58	4.8	2.29		
HEMBA1001734	2.37	2.38	4.28	4.78	6.06	4.27		
HEMBA1001736	2.3	2.12	2.87	3.5	4.69	5.33	*	+
HEMBA1001741	1.69	2.19	3.5	4.02	4.74	2.9		
HEMBA1001744	0.86	0.94	2.81	2	3.64	2.64		
HEMBA1001745	0.95	1.56	2.3	2.53	5.28	2.58		
HEMBA1001746	4.02	3.91	8.66	9.21	12.06	7.01		
HEMBA1001761	2.2	2.01	4.69	2.58	3.95	3.37		
HEMBA1001762	1.41	1.6	3.57	1.93	4.38	2.03		

(1029)

特開 2002-191363

2055

2056

HEMBA1001781	1.56	1.5	4.17	2.15	5.87	3.25		
HEMBA1001784	1.36	1.39	4.5	4.02	3.58	3.75		
HEMBA1001791	2.16	1.74	6.97	6.04	7.62	5.68		
HEMBA1001794	2.15	4.31	12.57	13.54	13.65	12.96		
HEMBA1001800	5.61	9.63	60.44	84.85	100.03	76.26	*	+
HEMBA1001803	2.84	4.25	5.36	4.27	7.02	3.67		
HEMBA1001804	6.2	8.13	20.95	29.84	26.86	24.39	*	+
HEMBA1001808	1.61	1.6	3.87	3.71	3.67	2.89		
HEMBA1001809	8.07	6.27	10.64	14.33	20.56	16.63	*	+
HEMBA1001811	8.32	7.83	16.8	22.75	21.75	17.6	*	+
HEMBA1001815	1.75	2.67	6.56	5.58	6.33	5.03		
HEMBA1001816	1.96	2.67	4.47	3.09	4.6	3.04		
HEMBA1001819	0.98	3.09	6.16	6.19	8.53	6.3		
HEMBA1001820	0.93	1.32	2.22	2.36	3.32	1.21		
HEMBA1001822	1.87	2.06	5.43	6.02	7.7	4.44		
HEMBA1001824	3.21	4.62	14.88	12.81	16.29	12.34		
HEMBA1001835	1.04	1.05	3.05	3.72	5.21	3.14		
HEMBA1001844	7.88	6.55	18.04	17.77	21.36	13.19		
HEMBA1001847	0.93	1.8	5.21	1.96	5.18	3.06		
HEMBA1001849	2.32	2.77	7.58	6.65	8.19	7.62		
HEMBA1001850	2.51	2.71	8.43	8.76	8.88	7.89		
HEMBA1001861	0.95	2.04	1.73	2.64	3.93	2.01		
HEMBA1001862	138.58	133.42	191.61	266.65	221.43	227.58	*	+
HEMBA1001864	1.31	1.16	2.44	4.79	2.88	2.59		
HEMBA1001866	1.49	2.39	7.45	7.67	7.07	4.61		
HEMBA1001869	7.55	6.84	10.82	10.31	7.69	9.02		
HEMBA1001871	29.48	30.98	54.77	63.07	62.43	66.59	*	+
HEMBA1001876	0.96	1.27	4.42	2.08	4.57	2.26		
HEMBA1001878	2.23	3.34	5.7	6.83	8.18	5.5		
HEMBA1001879	1.89	2.57	5.58	5.99	7.24	5.72		
HEMBA1001884	6.21	6.49	17.14	11.31	12.71	10.87		
HEMBA1001886	2.12	2.21	4.38	4.57	5.67	4.23		
HEMBA1001888	2	2.12	6.6	7.41	10.17	9.46	*	+
HEMBA1001890	4.03	3.67	7.6	6.8	7.01	4.4		
HEMBA1001896	1.34	1.61	2.62	2.27	4.12	3.01		
HEMBA1001899	33.43	39.48	61.77	106.52	41.01	101.91		
HEMBA1001904	76.64	122.45	233.99	299.33	174.47	322.82		
HEMBA1001910	1.4	1.93	3.23	2.53	6.35	2.97		
HEMBA1001911	8.36	8.75	10.86	21.15	16.9	13.23	*	+
HEMBA1001912	8.92	7.97	33.97	57	51.9	48.59	*	+
HEMBA1001913	4.89	6.19	17.29	18.56	14.16	16.85		
HEMBA1001915	1.35	2.61	4.49	3.3	5.63	2.46		
HEMBA1001918	15.23	13.29	21.07	17.07	14.31	12.13		
HEMBA1001921	4	3.5	4.38	5.2	5.35	4.86	*	+
HEMBA1001931	1.19	1.95	2.53	2.14	5.17	2.19		
HEMBA1001939	1.92	1.77	4.72	1.97	5.21	2.57		
HEMBA1001940	2.61	2.99	7.14	3.51	5.86	3.24		
HEMBA1001942	1.18	1.88	3.71	2.33	5.14	1.56		
HEMBA1001944	4.35	5.83	42.16	51.42	66.43	59.75	*	+
HEMBA1001945	0.98	2.3	2.95	2.98	3.4	2.21		

(1030)

特開2002-191363

2057

2058

HEMBA1001950	2.56	2.84	7.87	5.72	5.23	3.68		
HEMBA1001951	10.37	11.26	15.33	24.16	18.26	22.94	*	+
HEMBA1001958	1.04	1.28	2.58	3.1	4.83	2.54		
HEMBA1001960	6.87	6.28	13.93	10.02	12.99	12.47		
HEMBA1001962	1.01	1.08	4.19	1.58	4.24	1.67		
HEMBA1001964	1.39	3.45	4.13	2.54	4.45	3.39		
HEMBA1001967	6.06	5.65	9.33	14.45	10.5	13.18	*	+
HEMBA1001979	0.7	2.67	3.31	2.04	3.46	2.4		
HEMBA1001987	1.96	3.92	7.99	6.19	8.35	7.22		
HEMBA1001991	1.61	3.59	9.06	5.06	8.7	7.44		
HEMBA1002003	4.86	4.71	14.56	15.86	16.03	22.9		
HEMBA1002005	2.62	3.39	7.82	4.16	7.48	4.76		
HEMBA1002008	2.64	3.51	7.78	6.07	10.15	6.37		
HEMBA1002018	1.86	2.37	4.23	3.32	5.47	3.87		
HEMBA1002022	0.52	2.3	2.5	2.83	3.53	2.82		
HEMBA1002029	43.82	40.22	73.75	89.27	96.12	122.81	*	+
HEMBA1002030	2.23	2.88	4.32	3.88	4.26	4.67		
HEMBA1002035	1.69	1.75	3.82	5.43	5.14	3.75		
HEMBA1002037	4.47	4.34	6.69	4.5	6.47	7.94		
HEMBA1002038	4.12	3.13	7.74	6.36	8.8	5.42		
HEMBA1002039	2.46	3.43	7.03	6.74	8.99	6.37		
HEMBA1002042	5.52	5.55	9.8	10.55	13.01	9.94		
HEMBA1002043	3.81	3.79	11.32	12.53	12.64	13.95		
HEMBA1002048	2.76	2.31	3.81	2.03	4.32	3.44		
HEMBA1002049	1.72	2.35	5.55	4.43	5.2	5.2		
HEMBA1002053	7.33	6.91	14.52	11.09	15.25	13.63		
HEMBA1002055	9.81	8.76	10.65	18.44	13.58	17.78	*	+
HEMBA1002056	2.24	2.62	4.26	5.46	8.06	5.67	*	+
HEMBA1002061	2.24	2.51	4.58	4.17	5.34	4.58		
HEMBA1002080	46.55	49.5	54.6	91.78	122.41	83.23	*	+
HEMBA1002084	0.71	1.43	2.36	3.25	3.66	2.64	*	+
HEMBA1002085	0.97	1.47	2.87	3.45	4.52	3.74	*	+
HEMBA1002092	1.79	1.56	3.94	4.01	5.53	4.15		
HEMBA1002098	1.51	1.82	4.12	3.2	5.11	2.83		
HEMBA1002100	9.07	8.18	22.37	25.95	23.04	29.67		
HEMBA1002101	18.26	17.64	27.49	23.44	27.16	23.74		
HEMBA1002102	2.65	1.98	5.99	4.58	7.06	6.99		
HEMBA1002105	6.79	6.2	22.13	24.47	40.8	25.31		
HEMBA1002107	57.97	37.86	84.35	155.21	136.27	136.87	**	+
HEMBA1002113	6.77	4.75	17.24	12.05	12.83	14.78		
HEMBA1002119	3.85	3.28	24.05	24.29	30.66	27.04		
HEMBA1002125	7.03	6.73	10.43	11.38	7.34	11.85		
HEMBA1002131	9.71	9.72	20.58	14.97	19.12	13.71		
HEMBA1002133	3.67	3.52	6.32	6.97	9.51	9.46	*	+
HEMBA1002139	0.75	1.07	3.35	1.52	4.37	1.47		
HEMBA1002141	1.67	1.36	3.23	2.99	3.92	2.14		
HEMBA1002144	2.33	2.44	6.11	5.28	5.68	6.57		
HEMBA1002147	8.84	9.55	17.93	36.22	21.92	21.88		
HEMBA1002150	38.34	38.68	51.42	19.74	26.62	15.85	*	-
HEMBA1002151	3.76	3.36	9.95	11.15	14.23	9.84		

(1031)

特開2002-191363

2059

2060

HEMBA1002153	0.57	1.74	3.36	3.39	5.64	2.96		
HEMBA1002156	0.8	1.74	2.33	1.94	4.28	1.26		
HEMBA1002160	2.16	3.17	5.7	6.08	7.26	6.18		
HEMBA1002161	2.13	2.9	6.99	13.53	11.79	9.76	*	+
HEMBA1002162	2.65	2.17	7.76	5.61	6.27	7.81		
HEMBA1002163	12.02	12.04	19.93	34.48	19.96	27.11		
HEMBA1002164	6.58	10.55	59.92	71.46	78.61	67.82		
HEMBA1002166	21.88	18.32	39.58	57.35	49.05	46.09	*	+
HEMBA1002167	0.89	2.89	3.89	4.96	5.45	3.98		
HEMBA1002173	3.24	3.83	6.22	7.97	7.11	6.28		
HEMBA1002177	1.31	1.78	3.31	5.68	4.97	2.98		
HEMBA1002178	6.91	10.17	14.77	23.33	23.58	17.49	*	+
HEMBA1002179	53.56	46.86	94.4	58.33	85.22	54.47		
HEMBA1002185	2.75	4.07	13.4	11.73	16.23	14.56		
HEMBA1002188	5.76	7.57	10.27	11.86	12.9	9.8		
HEMBA1002189	1.98	2.85	4.96	5.23	4.63	4.71		
HEMBA1002191	0.67	2.16	4.96	3.47	5.44	2.81		
HEMBA1002192	2.98	2.83	4.91	7.53	8.35	4.57		
HEMBA1002195	2.96	3.27	6.6	10.35	10.11	7.27	*	+
HEMBA1002196	3.34	4.33	8.55	8.62	8.85	8.39		
HEMBA1002199	1.33	1.86	4.9	4.62	5.71	3.52		
HEMBA1002204	1.31	1.97	4.08	5.48	11.37	3.73		
HEMBA1002208	24.58	26.61	45.85	49.77	25.48	39.6		
HEMBA1002212	3.73	5.95	9.01	8.9	11.85	17.18		
HEMBA1002215	1.95	2.63	4.27	5.1	3.54	3.78		
HEMBA1002217	15.61	16.71	59.91	78.46	82.88	80.94	*	+
HEMBA1002220	1.11	2.07	4.1	3.58	3.39	2.33		
HEMBA1002226	2.17	3.13	9.18	10.47	12.61	9.58		
HEMBA1002227	39.9	47.13	92.5	109.42	65.74	71.79		
HEMBA1002229	4.5	4.77	13.39	11.16	13.55	12.49		
HEMBA1002237	1.73	3.22	4.08	3.71	5.64	3.41		
HEMBA1002239	9.36	13.83	72.18	100.62	109.3	113.84	*	+
HEMBA1002241	7	7.54	38.36	64.27	68.93	68.72	**	+
HEMBA1002253	1.11	2.44	3.33	2.3	4.42	2.68		
HEMBA1002257	1.83	2.65	4.11	3.18	3.6	1.74		
HEMBA1002259	1.12	2.17	2.69	3.12	3.6	2.67		
HEMBA1002262	6.95	7.37	19.16	14.43	14.78	17.04		
HEMBA1002265	1.35	1.63	3.7	3.75	6.23	2.43		
HEMBA1002267	16.87	20.81	22.76	32.99	16.96	27.5		
HEMBA1002270	3.73	4.79	7.49	8.18	13.43	8.7		
HEMBA1002286	1.03	1.86	5.42	2.85	5.53	0.98		
HEMBA1002290	4.73	3.7	7.52	5.16	8.88	4.38		
HEMBA1002302	6.12	9.63	45.04	45.66	49.69	45.23		
HEMBA1002304	3.28	3.42	6.88	5.57	6.97	4.34		
HEMBA1002307	45.71	53.69	92.31	71.87	55.79	61.03		
HEMBA1002316	2.16	3.29	4.63	3.04	5.32	2.41		
HEMBA1002319	1.97	2.96	4.46	8.32	9.58	8	**	+
HEMBA1002320	1.99	1.76	3.22	2.51	4.45	1.76		
HEMBA1002321	1.22	2.04	2.84	2.71	4.44	2.77		
HEMBA1002328	2.44	3.04	4.89	2.43	5.73	2.94		

(1032)

特開2002-191363

2061

2062

HEMBA1002333	3.88	4.27	7.14	7.37	10.84	7.5		
HEMBA1002337	3.02	3.62	5.5	5.98	7.49	7.48	*	+
HEMBA1002339	15.86	13.92	111.66	135.44	169.95	156.76	*	+
HEMBA1002341	0.8	2.08	3.22	2.71	4.34	2.35		
HEMBA1002348	2.84	2.78	7.14	3.69	6.73	4.49		
HEMBA1002349	1.28	1.44	3.59	2.24	4.64	2.59		
HEMBA1002353	1.83	3.04	4.03	4.61	7.72	5.68		
HEMBA1002356	6.05	6.96	17.53	14.27	16.02	16.1		
HEMBA1002357	114.85	156.08	306.32	300.67	286.5	328.19		
HEMBA1002360	7.18	8.32	8.29	14.57	14.46	13.78	**	+
HEMBA1002363	2.79	3.35	4.84	7.02	8.02	8.72	**	+
HEMBA1002365	1.7	2.7	2.7	1.63	3.12	2.67		
HEMBA1002370	1.43	1.78	2.37	1.53	4.2	1.9		
HEMBA1002374	4.55	4.53	7.79	8.33	10.27	9.11	*	+
HEMBA1002376	46.59	33.18	118.8	101.1	189.18	114.36		
HEMBA1002377	18.02	20.98	25.61	32.58	34.44	32.19	**	+
HEMBA1002380	5.68	6.36	16.28	17.43	21.85	18.83		
HEMBA1002381	1.52	1.8	4.16	4.12	7.16	4.94		
HEMBA1002384	1.79	3.09	3.69	5.67	4.27	5.71	*	+
HEMBA1002389	1.93	2.93	2.88	3.63	5.31	4.7	*	+
HEMBA1002396	21.16	20.01	36.93	14.29	14.94	19.1		
HEMBA1002402	125.09	124.52	168.42	100.85	107.79	164.62		
HEMBA1002417	1.41	1.07	4.27	2.17	3.19	2.6		
HEMBA1002419	1.42	2.38	3.8	1.81	4.59	2.41		
HEMBA1002420	9.55	11.97	14.11	16.34	18.28	16.16	*	+
HEMBA1002421	7.47	10.35	12.5	8.97	10.24	9.6		
HEMBA1002423	2.89	1.3	4.28	3.35	4.29	3.82		
HEMBA1002424	11.91	10.05	25.13	9.11	9.2	10.43		
HEMBA1002426	3.42	3.69	5.56	7.57	8.24	5.6	*	+
HEMBA1002430	0.39	1.41	2.51	1.7	4.19	2.85		
HEMBA1002439	1.59	1.94	4.17	2.69	4.46	6.8		
HEMBA1002441	31.85	29.77	27.79	48.74	52.7	34.64		
HEMBA1002454	0.62	1.48	2.27	1.76	2.43	1.55		
HEMBA1002458	3.17	5	8.09	10.24	10.55	11.71	*	+
HEMBA1002460	2.14	1.59	3.89	5.16	4.63	5.01	*	+
HEMBA1002462	5.18	3.83	9.52	11.92	9.29	8.58		
HEMBA1002465	0.93	1.96	2.26	1.46	3.13	1.27		
HEMBA1002469	6.88	7.27	45.87	49.02	75.59	67.74	*	+
HEMBA1002475	1.54	2.35	8.01	4.88	7.87	7.66		
HEMBA1002477	1.75	1.59	5.25	3.19	6.99	3.55		
HEMBA1002480	4.46	3.98	8.49	8.76	13	10.54		
HEMBA1002481	1.9	2.02	4.22	2.71	6.25	4.7		
HEMBA1002486	3.62	3.98	10.08	9.98	8.04	9.75		
HEMBA1002490	2.02	3.08	5.7	8.76	9.64	7.65	*	+
HEMBA1002495	2.37	2.29	3.78	3.92	4.79	4.08		
HEMBA1002498	0.95	2.14	2.97	1.83	5.09	2.14		
HEMBA1002501	2.96	4.73	14.13	19.98	23.55	17.54	*	+
HEMBA1002503	1.7	2.52	5.11	4.68	7.06	2.97		
HEMBA1002504	1.95	2.19	5.99	6.68	7.09	4.65		
HEMBA1002508	1.48	2.59	5.99	7.8	7.47	5.65		

(1033)

特開2002-191363

2063

2064

HEMBA1002513	1.31	1.7	4.85	3.91	7.67	3.02		
HEMBA1002515	1.17	1.82	3.04	2.67	5.1	2.89		
HEMBA1002524	1.67	2.09	2.53	4.44	4.49	3.82	**	+
HEMBA1002538	4.68	4.14	7.39	9.31	8.91	7.86	*	+
HEMBA1002542	3.31	3.27	6.77	10.11	9.3	7.74	*	+
HEMBA1002544	1.42	2.24	3.33	2.69	6.59	3.24		
HEMBA1002546	31.01	31.64	56.69	95.52	83.15	72.77	*	+
HEMBA1002547	3.13	3.22	8.44	20.11	20.37	17.21	**	+
HEMBA1002550	5.46	3.86	10.87	10.85	11.2	10.23		
HEMBA1002551	2.15	3.09	5.8	3.7	5.08	3.08		
HEMBA1002552	2.21	2.06	8.39	5.66	6.55	5.68		
HEMBA1002555	1.54	1.78	4.56	2.27	4.4	2.97		
HEMBA1002558	2.74	3.26	7.02	8.08	7.47	8.27		
HEMBA1002561	1.01	1.58	5.26	4.42	5.08	3.87		
HEMBA1002562	0.59	0.83	2.34	3.29	3.29	2.36		
HEMBA1002568	1.71	1.16	3.09	3.06	3.26	3.6		
HEMBA1002569	3.8	4.67	10.32	7.29	8.59	5.14		
HEMBA1002570	5.22	4.72	9.84	6.07	10.29	12.99		
HEMBA1002574	24.62	22.75	26.01	44.47	30.74	40.85	*	+
HEMBA1002583	4.07	4.52	8.07	6.64	6.43	8.47		
HEMBA1002587	9.78	10.9	19.23	24.67	18.08	20.4		
HEMBA1002590	2.51	2.58	7.47	5.35	5.6	4.05		
HEMBA1002592	2.51	3.03	6.4	6.34	7.84	5.1		
HEMBA1002595	1.66	2.13	3.1	4.12	4.25	2.68		
HEMBA1002609	4.47	6.27	51	68.51	85.44	66.33	*	+
HEMBA1002617	6.31	4.76	7.99	7.25	6.84	6.48		
HEMBA1002619	3.33	4.8	5.99	7.86	6.14	7.27		
HEMBA1002621	1.21	2.94	3.09	3.24	4.17	2.03		
HEMBA1002624	4.6	5.19	19.48	22.04	24.39	26.87	*	+
HEMBA1002628	3.37	3.64	6.41	6.08	6.11	4.1		
HEMBA1002629	2.71	2.24	4.66	3.77	7.98	4.48		
HEMBA1002632	1.39	2.23	5.16	4.29	5.49	4.58		
HEMBA1002645	1.77	1.98	6.43	4.68	6.91	5.37		
HEMBA1002651	1.87	2.73	4.73	4.68	4.83	3.74		
HEMBA1002652	3.38	5.27	6.21	6.09	8.66	7.92		
HEMBA1002659	2.84	3.86	4.8	6.32	8.18	9.6	*	+
HEMBA1002661	3	2.71	6.19	4.41	6.93	4.93		
HEMBA1002666	1.74	2.47	4.21	2.95	4.25	1.41		
HEMBA1002667	1.39	2.25	3.91	2.79	5.24	1.94		
HEMBA1002673	16.08	19.36	30.31	32.54	35.18	29.96		
HEMBA1002678	2.11	2.33	7.44	5.39	5.98	4.22		
HEMBA1002679	1.23	2.33	5.25	3.7	7.48	3.81		
HEMBA1002688	1.74	2.98	8.3	8.33	11.41	7.86		
HEMBA1002696	1.7	2.79	2.92	3.48	6.13	3.32		
HEMBA1002703	2.95	3.88	10.15	8.35	9.73	9.21		
HEMBA1002706	4.97	4.24	8.99	5.07	7.16	5.54		
HEMBA1002712	2.39	3.94	8.67	8.4	10.9	10.57		
HEMBA1002715	7.92	9.81	49.65	79.65	93.63	79.61	*	+
HEMBA1002716	3.93	4.26	5.53	4.63	5.02	4.53		
HEMBA1002718	11.79	12.87	17.77	24.16	18.07	24.3	*	+

(1034)

特開2002-191363

2065

2066

HEMBA1002728	2.37	3.1	5.01	5.52	5.94	4.42		
HEMBA1002730	1.13	2.48	5.86	3.71	6.19	4.61		
HEMBA1002734	2.89	3.54	8.82	8.6	10.7	10.59		
HEMBA1002742	1.94	2.06	3.96	1.86	4.27	2.74		
HEMBA1002746	1.2	2.86	4.61	2.83	4.43	2.94		
HEMBA1002748	2.19	1.75	4.01	5.36	5.98	3.92		
HEMBA1002750	1.99	2.46	3.45	6.74	6.39	6.27	**	+
HEMBA1002755	1.85	3.1	5.31	5.96	6.62	5.16		
HEMBA1002759	1.93	3.12	7.98	4.65	7.92	7.08		
HEMBA1002763	9.62	12.05	74.52	68.84	88.82	77.22		
HEMBA1002767	4.48	5.85	5.8	8.88	6	6.13		
HEMBA1002768	2.99	3.76	6.2	3.46	8.3	3.04		
HEMBA1002769	1.47	2.35	2.82	3.46	5.21	3.49		
HEMBA1002770	5.89	5.83	12.41	14.24	22.53	15.53	*	+
HEMBA1002777	1.6	1.9	2.58	2.29	4.74	3.76		
HEMBA1002779	10.92	7.6	16	17.39	19.81	19.36	*	+
HEMBA1002780	2.6	2.77	6.82	6.43	6.89	6.35		
HEMBA1002790	3.14	2.52	10.6	7.26	8.67	9.25		
HEMBA1002794	1.52	2.28	5.49	3.68	6.8	4.45		
HEMBA1002798	1.33	1.59	3.61	2.77	5.12	4		
HEMBA1002801	2.13	2.25	3.64	3.12	6.93	5.24		
HEMBA1002810	4.56	3.99	7.85	10.27	17.31	11.1	*	+
HEMBA1002816	2.24	1.97	2.88	5.34	5.05	4.8	**	+
HEMBA1002818	24.6	23.26	95.11	130.84	121.74	135.78	*	+
HEMBA1002820	1.95	2.63	6.41	6.96	6.99	6.04		
HEMBA1002826	1.96	1.48	2.99	3.21	4.84	3.59		
HEMBA1002833	8.71	7.46	19.84	20.18	21.16	20.04		
HEMBA1002850	1.16	1.94	3.67	3.87	4.96	5.11	*	+
HEMBA1002862	9.06	9.31	17.9	20.11	25.3	13.43		
HEMBA1002863	2.47	2.93	5.28	6.16	8.44	6.52	*	+
HEMBA1002867	1.51	1.17	2.4	2.3	3.28	1.87		
HEMBA1002876	3.9	3.54	5.48	5.61	5.78	6.48		
HEMBA1002886	1.28	1.56	2.45	1.83	3.13	2.71		
HEMBA1002896	5.82	3.82	9.38	7.22	11.23	8.51		
HEMBA1002913	2.37	2.22	4.56	4.19	4.28	3.11		
HEMBA1002921	0.97	0.81	2.36	1.82	2.41	1.41		
HEMBA1002924	1.07	1.2	2.86	2.11	4.41	3.27		
HEMBA1002934	6.01	5.17	10.48	9.93	15.27	13.16		
HEMBA1002935	4.27	2.55	6.59	7.1	5.34	7.14		
HEMBA1002937	4.61	5.71	9.4	10.82	8.36	7.36		
HEMBA1002939	2.21	2.92	5.39	5.51	5.7	3.26		
HEMBA1002944	1.45	1.97	4.66	3.1	5.68	3.21		
HEMBA1002951	5.88	7.88	10.99	6.04	12.17	5.67		
HEMBA1002954	2.4	4.57	6.12	6.09	7.78	4.78		
HEMBA1002962	3.93	6.02	9.14	13.42	15.92	12.44	*	+
HEMBA1002968	1.22	1.71	4.32	5.34	4.07	5.3		
HEMBA1002970	1.13	1.13	3.14	2.5	3.72	3.13		
HEMBA1002971	0.96	2.02	2.75	2.02	3.71	2.43		
HEMBA1002973	1.68	3.36	7.84	6.19	10.81	6.31		
HEMBA1002978	2.09	3.81	4.35	5.49	5.3	4.22		

(1035)

特開2002-191363

2067

2068

HEMBA1002981	1.82	2.51	4.01	9.33	7.48	6.53	**	+
HEMBA1002985	0.83	1.92	4.91	4.74	5.59	4.13		
HEMBA1002986	2.72	4.88	6.67	14.7	14.62	13	**	+
HEMBA1002988	1.77	2.36	4.25	4.2	5.67	3.46		
HEMBA1002992	8.73	11.38	68.65	83.81	96.4	94.02	*	+
HEMBA1002995	6.13	6.97	11.94	8.64	11.47	14.09		
HEMBA1002997	5.77	6.33	9.6	12.88	10.65	8.75		
HEMBA1002999	1.36	2.77	2.84	2.48	3.92	3.31		
HEMBA1003004	0.78	1.39	1.96	1.86	3.32	1.37		
HEMBA1003006	2.03	1.84	4.26	5.44	8.08	5.87	*	+
HEMBA1003008	1.58	1.26	2.83	3.4	5.4	2.28		
HEMBA1003021	1.72	2.09	5.98	8.49	8.58	6.67	*	+
HEMBA1003027	1.79	1.73	4.47	2.11	6.17	3.72		
HEMBA1003029	16.39	17.36	46.06	37.07	42.91	45.58		
HEMBA1003031	33.04	32.41	50.08	48.18	24.56	40.3		
HEMBA1003032	3.42	6.52	7.98	8.81	6.53	9.45		
HEMBA1003033	2.36	4.11	6.85	7.85	8.94	9	*	+
HEMBA1003034	2.43	3.17	7.63	8.24	7.47	8.23		
HEMBA1003035	1.24	2	2.59	2.88	3.46	1.93		
HEMBA1003037	1.74	2.09	6.21	4.13	7.36	3.43		
HEMBA1003041	3.4	4.14	8.51	10.28	10.48	9		
HEMBA1003046	11.44	11.53	28.31	33.77	21.19	36.32		
HEMBA1003047	2.02	2.35	5.11	4.57	5.41	4.03		
HEMBA1003048	1.8	2.96	3.76	7.97	7.47	9.3	**	+
HEMBA1003064	3.7	4.12	15.74	15.78	25.09	19.36		
HEMBA1003067	1.92	2.31	7.09	4.56	7.96	3.23		
HEMBA1003071	5.24	5	8.74	11.32	10.02	8.7		
HEMBA1003072	2.81	3.22	5.7	4.43	3.65	5.17		
HEMBA1003076	20.6	21.34	31.6	41.86	28.3	32.74		
HEMBA1003077	1.41	1.58	4.37	1.68	4.03	2.27		
HEMBA1003078	2.02	1.92	2.4	3.14	4.9	3.38	*	+
HEMBA1003079	2.72	2.66	6.42	5.88	7.13	4.48		
HEMBA1003083	1.56	2.11	3.94	4.42	6.23	4.59		
HEMBA1003086	2.5	2.72	5.27	4.09	5.78	4.68		
HEMBA1003090	5.14	4.79	13.3	11.57	12.88	12.73		
HEMBA1003094	0.82	1.67	2.94	2.51	3.22	2.14		
HEMBA1003096	8.6	8.76	15.55	10.1	13.7	10.97		
HEMBA1003098	3.88	5.66	7.38	9.42	7.11	8.4		
HEMBA1003101	4.73	5.48	7.29	9.04	6.59	6.36		
HEMBA1003109	2.88	3.42	4.72	5.73	6.93	7.22	*	+
HEMBA1003114	2.87	4.67	5.67	6.47	7.94	5.69		
HEMBA1003117	2.1	3.41	4.4	3.36	4.99	2.44		
HEMBA1003120	3.02	2.65	5.55	3.23	7.38	4.29		
HEMBA1003129	2.47	2.6	6.66	10.28	6.19	7.28		
HEMBA1003133	2.05	4.74	7.61	7.74	7.59	5.59		
HEMBA1003136	2.64	3.59	5.25	5.37	5.88	4.49		
HEMBA1003142	2.01	2.27	6.15	6.62	6.35	5.34		
HEMBA1003148	1.3	1.4	2.82	1.49	4.26	1.6		
HEMBA1003151	1.91	2.08	4.23	2.9	5.34	4.24		
HEMBA1003152	3.27	1.98	5.84	4.74	5.71	2.58		

(1036)

特開 2002-191363

2069

2070

HEMBA1003157	1.23	1.88	2.58	4.2	5.38	3.21	*	+
HEMBA1003166	6.14	6.06	14.06	22.98	18.03	21.74	*	+
HEMBA1003171	1.3	2.28	2.23	2.62	3.09	2.53		
HEMBA1003175	1.54	2.63	4.2	3.54	4.52	4.11		
HEMBA1003179	4.66	5.95	37.4	36.91	43.86	45.13		
HEMBA1003186	2.58	3.17	7.13	6.71	6.71	5.78		
HEMBA1003196	3.04	3.79	7.33	6.95	8.31	5.18		
HEMBA1003197	0.46	1.51	2.86	1.85	3.97	1.09		
HEMBA1003199	1.26	1	2.32	1.66	3.22	2.47		
HEMBA1003202	2.86	3.49	5.69	9.44	10.48	11.14	**	+
HEMBA1003204	1.67	2.46	3.35	4.99	4.72	4.81	**	+
HEMBA1003210	6.48	7.36	11.66	12.02	12.1	14.78		
HEMBA1003212	1.4	2.87	5.52	7.58	8	5.7		
HEMBA1003218	1.2	1.26	1.71	1.24	4.35	1.36		
HEMBA1003220	34.65	32.6	73.43	78.35	79.82	83.89		
HEMBA1003222	2.37	3.03	3.41	3.04	6.13	4.29		
HEMBA1003225	1.95	2.07	3.34	1.59	3.45	2.05		
HEMBA1003229	2.37	1.91	2.4	5.62	5.1	4.9	**	+
HEMBA1003230	7.83	7.14	12.08	11.08	11.44	10.09		
HEMBA1003235	0.91	1.33	4.32	4.98	5.25	5.44	*	+
HEMBA1003236	5.54	5.43	10.62	11.5	15.4	13.97	*	+
HEMBA1003250	1.41	1.4	2.68	1.76	2.98	2.42		
HEMBA1003252	4.96	7.17	16.59	17.06	18.68	14.37		
HEMBA1003257	2.7	3.33	7.33	8.25	8.83	6.78		
HEMBA1003268	0.95	0.44	1.92	1.92	3.36	1.81		
HEMBA1003273	1.4	1.38	2.96	2.5	3.37	5.1		
HEMBA1003276	1.13	1.99	3.18	4.21	4.42	3.98	*	+
HEMBA1003277	0.95	0.83	1.85	0.56	1.63	1.34		
HEMBA1003278	1.07	1.18	3.49	1	4.56	2.32		
HEMBA1003280	2.37	2.6	4.59	3.08	4.91	4.2		
HEMBA1003281	1.83	1.29	3.53	1.79	3.85	2.48		
HEMBA1003284	1.24	1.91	3.43	3.03	5.05	3.32		
HEMBA1003286	7.75	6.73	34.23	45.25	71.61	51.97	*	+
HEMBA1003291	1.65	1.91	4.84	3.21	3.32	3.25		
HEMBA1003294	1.89	3.5	7.47	4.86	5.7	5.62		
HEMBA1003296	4.74	8.32	46.61	63.82	70.23	54.45	*	+
HEMBA1003304	0.77	1.44	2.88	2.91	5.37	1.87		
HEMBA1003306	4.37	6.3	10.28	15.7	17.76	11.56	*	+
HEMBA1003309	0.91	1.9	2.85	2.87	4	2.3		
HEMBA1003314	1.43	2.26	3.82	4.48	3.52	4.02		
HEMBA1003315	6.37	4.38	10.14	15.2	16.23	17.88	**	+
HEMBA1003322	4.81	5.92	10.9	8.46	10.83	8.07		
HEMBA1003326	1.94	3.97	5.55	2.93	7.4	3.68		
HEMBA1003327	0.81	1.61	3.63	2.36	4.3	2.28		
HEMBA1003328	0.76	2.43	5.38	4.25	5.51	5.06		
HEMBA1003330	2.27	2.81	4.84	4.66	5.83	6.94		
HEMBA1003348	3.22	2.45	11.3	11.28	13.98	16.37		
HEMBA1003369	2.39	2.6	7	9.64	8.65	5.33		
HEMBA1003370	3.14	3.6	8.85	12.54	10.83	13.98	*	+
HEMBA1003373	1.12	1.3	3.4	2.14	5.05	2.94		

(1037)

特開2002-191363

2071

2072

HEMBA1003376	3.75	2.83	7.71	9.83	12.46	10.39	*	+
HEMBA1003380	1.12	2.3	3.63	2.25	3.9	2.57		
HEMBA1003384	0.98	1.71	2.91	2.11	4.78	2.14		
HEMBA1003387	1.3	1.24	2.14	1.83	3.24	1.98		
HEMBA1003392	2.51	2.28	3.43	5.21	5.91	4.44	*	+
HEMBA1003395	1.02	1.45	2.84	4.06	4.29	2.18		
HEMBA1003399	1.03	1.4	3.27	3.21	3.26	2.19		
HEMBA1003400	1.36	2.22	4.64	3.23	7.19	5.22		
HEMBA1003402	1.62	1.74	3.29	2.32	4.22	2.59		
HEMBA1003403	7.13	9.32	50.9	66.1	66.81	77.49	*	+
HEMBA1003408	3.68	4.5	7.27	6.02	5.77	7.71		
HEMBA1003412	5.08	6.79	8.35	10.96	8.79	9.75		
HEMBA1003417	5.71	6.5	10.15	8.18	8.86	7.36		
HEMBA1003418	4.01	5.12	6.53	7.37	11.45	9.3	*	+
HEMBA1003420	16.29	17.91	35.46	33.32	34.37	32.89		
HEMBA1003425	0.76	1.65	3.06	2.33	3.58	2.21		
HEMBA1003433	1.4	2.43	3.34	4.88	4.54	4.09	*	+
HEMBA1003440	11.39	12.08	19.86	24.13	13.99	24.26		
HEMBA1003442	4.37	4.67	4.94	3.54	6.73	5.96		
HEMBA1003447	7.55	9.08	49.72	65.41	63.46	65.15	*	+
HEMBA1003453	21.03	22.03	42.15	27.85	29.02	27.64		
HEMBA1003461	1.5	2.13	3.49	2.6	3.63	2.2		
HEMBA1003463	2.82	3.68	6.02	5.97	3.84	6.41		
HEMBA1003465	1.77	2.21	6.31	4.75	5.02	3.82		
HEMBA1003480	2.58	3.91	8.62	9.63	9.6	9.42		
HEMBA1003485	7.06	4.84	5.29	6.13	7.26	5.52		
HEMBA1003487	1.8	1.85	3.4	7.12	6.39	6.79	**	+
HEMBA1003492	1.42	1.95	4.11	2.41	5.87	2.1		
HEMBA1003494	9.36	8.61	12.16	18.24	18.69	17.83	**	+
HEMBA1003497	2.19	2.16	3.29	3.35	6.06	2.97		
HEMBA1003503	0.98	1.74	3.37	5.04	3.18	2.13		
HEMBA1003511	0.99	2.19	3.7	2.3	4.42	2.5		
HEMBA1003528	3.33	4	6.51	5.77	5.04	4.46		
HEMBA1003530	1.33	0.85	3.62	1.97	3.15	2.45		
HEMBA1003531	1.14	1.72	5.39	4.74	7.24	4.51		
HEMBA1003532	12.97	14.66	34.3	28.69	25.31	31.26		
HEMBA1003538	2.54	2.4	17.88	14.54	21.58	16.83		
HEMBA1003545	0.68	2.08	3.17	1.85	3.6	2.17		
HEMBA1003546	1.27	2.03	1.68	1.98	2.15	2.42		
HEMBA1003548	1.4	3.18	3.6	1.41	4.15	2.23		
HEMBA1003553	31.29	31.45	47.99	54.36	41.34	45.65		
HEMBA1003555	1.39	2.73	4.81	3.53	4.48	5.19		
HEMBA1003556	1.24	1.76	2.96	3.14	5.75	3.31		
HEMBA1003560	1.89	2.66	7.87	10.08	13.24	9.9	*	+
HEMBA1003565	54.27	66.88	96.28	121.29	139.88	148.68	*	+
HEMBA1003568	1.86	2.27	3.24	2.36	7.41	2.78		
HEMBA1003569	2.93	2.61	2.96	5.07	3.95	4.53	**	+
HEMBA1003571	3.53	2.33	3.8	5.19	5.3	5.83	*	+
HEMBA1003579	3.51	4.29	4.83	3.79	5.68	5.91		
HEMBA1003580	3.82	4.09	4.96	3.11	4.41	3.53		

(1038)

特開2002-191363

2073

2074

HEMBA1003581	0.82	2.62	2.07	1.63	3.19	2.4		
HEMBA1003591	10.8	11.44	30.24	33.74	35.7	36.88	*	+
HEMBA1003595	0.93	1.16	2.46	2.98	4.02	2.01		
HEMBA1003597	3.15	3.18	8.74	10.82	11.39	11.59	*	+
HEMBA1003598	0.58	0.93	1.33	2.62	1.83	1.61	*	+
HEMBA1003600	3.71	4.19	13.35	14.77	13.86	16.69		
HEMBA1003602	2.84	2.64	4.89	5.89	6.97	9.14	*	+
HEMBA1003604	2.3	3.35	5.67	6.63	8.29	8.16	*	+
HEMBA1003610	2.33	3.2	4.48	6.12	5.64	6.81	*	+
HEMBA1003615	1.76	2.61	5.23	4.95	5.21	4.96		
HEMBA1003617	3.59	3.54	8.59	6.92	11.37	8.5		
HEMBA1003620	5.76	6.01	4.98	13.48	17.69	12.58	**	+
HEMBA1003621	1.6	1.66	3.19	4.52	5.42	5.08	**	+
HEMBA1003622	0.96	0.69	1.38	1.47	3.17	2.25		
HEMBA1003630	0.78	1.02	1.95	1.68	2.97	1.55		
HEMBA1003637	0.66	1.93	2.59	2.11	3.11	2.63		
HEMBA1003640	2.33	2.1	5.27	4.16	5.68	5.5		
HEMBA1003645	1.12	1.2	4.41	2.3	3.82	3.06		
HEMBA1003646	0.94	1.21	1.76	1.25	3.25	1.8		
HEMBA1003647	0.49	2.15	3.27	2.46	3.79	2.21		
HEMBA1003656	3.32	3.77	6.96	17.01	10.45	13.78	*	+
HEMBA1003662	1.37	2.08	1.54	5.2	3.81	4.91	**	+
HEMBA1003666	23.84	17.7	51.57	21.97	21.85	24.71		
HEMBA1003667	4.74	3.63	6.03	4.61	6.22	7.09		
HEMBA1003670	0.83	0.65	1.94	1.18	2.61	1.51		
HEMBA1003674	32.16	29.41	63.99	118.95	138.25	123.17	**	+
HEMBA1003677	1.84	2.06	4.28	2.32	5.31	3.78		
HEMBA1003679	1.2	1.68	3.72	2.22	6.19	3.23		
HEMBA1003680	4.55	4.68	20.52	27.26	28.13	28.07	*	+
HEMBA1003684	1.57	1.9	3.98	4	3.65	4.47		
HEMBA1003690	6.22	7.41	8.65	7.94	9.93	7.33		
HEMBA1003692	2.41	3.82	7.23	8	8.28	7.7		
HEMBA1003702	2.64	3.82	4.83	7.11	6.86	6.07	*	+
HEMBA1003711	1.06	1.21	3.39	2.93	3.88	2.37		
HEMBA1003714	1.31	1.26	2.13	1.61	2.45	1.42		
HEMBA1003715	1.46	2.7	6.58	10.21	9.15	6.87	*	+
HEMBA1003717	1.91	2.31	3.91	3.03	3.66	4.38		
HEMBA1003720	0.81	2.6	5.07	4.16	4.16	4.21		
HEMBA1003725	0.83	1.57	2.47	3.22	4.91	3.17	*	+
HEMBA1003728	1.28	2.48	3.4	2.65	4.36	2.72		
HEMBA1003729	0.98	2.35	2.85	3.6	4.36	3.52	*	+
HEMBA1003732	1.11	1.52	3.49	3.01	2.75	1.88		
HEMBA1003733	1.18	1.9	2.94	3.7	4.95	3.92	*	+
HEMBA1003742	5.15	7.3	5.95	21.53	22.58	19.56	**	+
HEMBA1003743	1.37	1.76	3.21	4.13	4.36	3.68	*	+
HEMBA1003758	3.26	3.29	11.72	10.07	16.03	12.24		
HEMBA1003760	0.82	2.43	3.09	1.92	4.19	3.16		
HEMBA1003764	0.88	2.06	4.9	1.86	4.36	4.24		
HEMBA1003769	6.61	8.95	15.57	22.58	17.05	20.77	*	+
HEMBA1003773	2.16	3.5	4.48	5.8	6.7	5.98	*	+

(1039)

特開2002-191363

2075

2076

HEMBA1003783	3.12	3.11	4.95	8.58	8.64	8.27	**	+
HEMBA1003784	0.46	1.37	2.89	2.37	2.45	1.77		
HEMBA1003794	3.48	3.61	13.64	11.98	11.61	15.55		
HEMBA1003799	1.09	1.05	4.29	2.44	5.06	3.76		
HEMBA1003803	7.58	6.67	12.05	11.68	7.64	8.41		
HEMBA1003804	1.08	2.53	4.45	1.86	3.35	2.05		
HEMBA1003805	6.41	7.87	10.45	15.74	7.09	11.93		
HEMBA1003807	1.52	1.53	3.21	2.71	6.32	2.37		
HEMBA1003810	1.72	3.29	6.06	5.98	4.85	4.55		
HEMBA1003827	2.71	4.55	12.08	13.28	10.48	15.35		
HEMBA1003836	3.42	4.84	10.27	11.16	12.81	9.96		
HEMBA1003838	16.58	16.15	31.32	34.24	33.25	35.46		
HEMBA1003843	4.6	6.54	7.01	13.61	6.48	11.42		
HEMBA1003846	19.54	21.94	61.32	72.86	70.58	83.4	*	+
HEMBA1003856	1.41	1.66	2.85	2.07	4.03	2.51		
HEMBA1003857	2.89	3.1	5.85	5.89	8.29	6.88		
HEMBA1003864	1.56	2.61	4.04	3.32	4.03	2.75		
HEMBA1003866	0.89	0.75	2.21	1.66	2.23	0.73		
HEMBA1003868	10.92	10.88	18.59	13.26	7.59	15.72		
HEMBA1003879	0.95	1.33	3.16	3.49	4.42	3.09		
HEMBA1003880	1.81	2.35	2.78	3.53	4.78	2.3		
HEMBA1003884	10.97	11.37	39.03	54.69	62.46	57.8	*	+
HEMBA1003885	4.59	4.82	7.14	9.19	6.32	8.41		
HEMBA1003887	3.58	4.93	7.7	8.65	7.93	8.18		
HEMBA1003890	4.2	4.48	7.18	7.53	9.1	6.26		
HEMBA1003893	4.38	6.39	9.53	8.75	13.24	9.94		
HEMBA1003896	4.15	4.15	10.62	7.4	9.12	6.43		
HEMBA1003902	1.39	3.78	5.09	4.91	6.42	5.1		
HEMBA1003904	0.87	2.16	2.46	2.82	4.32	2.11		
HEMBA1003908	1.18	1.3	2.89	2.12	5.25	1.43		
HEMBA1003926	14.46	12.2	39.79	45.5	34.97	55.56		
HEMBA1003937	2.75	3.31	5.38	4.3	6.85	4.57		
HEMBA1003939	2.43	2.48	6.56	8.3	13.32	8.04		
HEMBA1003940	2.45	3.08	5.01	4.29	6.22	5.55		
HEMBA1003941	1.4	2.26	2.48	3.37	4.57	4.42	*	+
HEMBA1003942	1.63	2.88	3.13	2.01	3.85	2.22		
HEMBA1003945	12.57	13.75	22.75	20.99	14.77	19.74		
HEMBA1003949	1.4	1.9	3.53	3.29	6.22	4.14		
HEMBA1003950	3.46	4.86	6.49	14.69	17.53	13.02	**	+
HEMBA1003953	1.91	1.6	5.14	0.72	3.97	1.44		
HEMBA1003958	5.16	3.6	7.47	7.54	9.45	6.64		
HEMBA1003959	2.42	2.72	5.72	5.5	5.5	9.02		
HEMBA1003960	3.25	5.81	34.7	24.04	26.4	28.28		
HEMBA1003966	9.63	8.28	16.73	16.75	17.67	19.84		
HEMBA1003967	1.75	3.06	3.47	3.48	3.6	3.27		
HEMBA1003968	0.97	2.14	2.55	2.49	4.56	1.82		
HEMBA1003974	634.2	699.64	821.36	986.23	1340.97	1248.21	*	+
HEMBA1003976	1.05	1.84	3.36	1.21	3.27	2.04		
HEMBA1003977	1.48	2.07	1.99	1.41	3.49	2.15		
HEMBA1003978	2.91	3.72	3.54	3.77	6.18	3.53		

(1040)

特開2002-191363

2077

2078

HEMBA1003981	9.01	6.77	14.06	12.05	11.49	18.27		
HEMBA1003982	102.64	103.61	302.15	380.08	375.9	466.69	*	+
HEMBA1003985	1.18	1.9	2.43	3.21	3.79	2.18		
HEMBA1003987	3.04	2.23	3.1	2.56	4.34	5.53		
HEMBA1003989	1.62	1.77	4.56	3.79	5.12	3.31		
HEMBA1004000	1.63	2.35	5.05	3.46	5.35	4.18		
HEMBA1004006	2.79	2.88	12.86	16.29	22.13	19.73	*	+
HEMBA1004007	0.7	1.92	5.28	3.03	5.18	4.72		
HEMBA1004010	67.4	61.25	98.24	112.56	96.78	136.86		
HEMBA1004011	0.48	1.74	2.18	2.58	3.29	1.62		
HEMBA1004012	0.79	1.84	2.3	3.11	4.8	3.53	*	+
HEMBA1004015	2.68	4.15	5.38	8.68	10.65	9.21	**	+
HEMBA1004024	1.47	2.73	5.65	5.68	8.26	8.07		
HEMBA1004029	1.93	3.1	3.03	4.6	8.38	13.11		
HEMBA1004038	1.04	1.24	1.55	1.18	3.38	1.59		
HEMBA1004042	0.89	1.42	2.22	1.58	4	2.48		
HEMBA1004045	0.28	0.94	2.42	3.07	3.32	2.53		
HEMBA1004048	4.16	4.16	12.1	19.93	14.84	22.3	*	+
HEMBA1004049	3.56	3.18	4.87	4.92	6.83	5.48		
HEMBA1004051	136.19	118.77	205.49	243.62	283.22	223.29	*	+
HEMBA1004053	5.11	4.64	8.92	25.25	27.24	21	**	+
HEMBA1004055	2.28	3.2	4.24	2.15	5.51	2.86		
HEMBA1004056	3.78	3.07	6.73	5.3	10.99	9.56		
HEMBA1004060	0.86	1	1.7	0.78	3.94	1.65		
HEMBA1004061	4.76	3.94	6.44	7.37	12.64	8.57		
HEMBA1004067	10.12	14.76	90.67	108.89	125.21	128.6	*	+
HEMBA1004071	7.51	7.77	16.52	17.31	12.23	13.37		
HEMBA1004074	0.78	1.93	3.97	4.48	7.06	5.69	*	+
HEMBA1004078	3.87	2.95	5.22	6.52	6.2	6.87	*	+
HEMBA1004085	1.05	1.19	2.83	3.57	4.57	2.45		
HEMBA1004086	3.38	4.95	6	8.92	8.09	6.51	*	+
HEMBA1004097	1.18	1.13	2.97	3.66	3.28	2.97		
HEMBA1004100	3.85	4.81	8.96	6.9	9.64	9.55		
HEMBA1004103	2	2.91	6.25	6.25	7.24	7.38		
HEMBA1004110	3	3.77	5.43	4.18	4.23	5.02		
HEMBA1004111	3.96	7.64	44.2	53.81	60.1	57.3	*	+
HEMBA1004124	7.14	10.51	60.12	83.27	97.96	83.59	*	+
HEMBA1004130	3.12	3.46	10.29	9.45	6.84	8.43		
HEMBA1004131	2.14	2.12	3.06	4.08	3.73	3.21	*	+
HEMBA1004132	0.77	2.22	4.84	3.94	6.31	4.2		
HEMBA1004133	0.69	1.77	2.56	3.28	5.17	3.22		
HEMBA1004138	0.89	1.19	3.05	2.21	4.11	1.83		
HEMBA1004143	7.1	7.48	17.43	18.83	15	17.6		
HEMBA1004146	0.89	2.03	3.01	2.96	4.21	2.69		
HEMBA1004148	1.85	1.57	2.13	2.25	3.38	1.99		
HEMBA1004149	1.54	1.44	2.77	2.83	2.59	3.32		
HEMBA1004150	0.49	1.06	2.15	2.31	1.58	1.08		
HEMBA1004154	2.24	1.64	5.28	6.28	7.07	4.61		
HEMBA1004164	1.84	2.23	5.63	6.89	7.13	5.81		
HEMBA1004168	2.16	2.24	4.69	3.9	5.32	7.84		

(1041)

特開 2002-191363

2079

2080

HEMBA1004199	1.37	1.92	2.34	3.17	3.66	1.8		
HEMBA1004200	0.84	1.98	3	1.5	4.05	1.78		
HEMBA1004201	4.87	5.68	17.64	26.94	32.17	25.65	*	+
HEMBA1004202	7.7	10.5	9.9	18.08	16.29	15.77	**	+
HEMBA1004203	1.63	2.31	3.66	4.5	5.3	4.44	*	+
HEMBA1004207	1.9	3.24	3.62	5.73	6.23	6.2	**	+
HEMBA1004210	1.13	1.72	2.67	1.95	4.14	1.87		
HEMBA1004225	1.1	2.47	5.23	5.96	7.12	5.4		
HEMBA1004227	2.17	4.44	3.86	5.14	5.71	5.16		
HEMBA1004235	2.68	2.91	3.74	5.79	5.78	4.44	*	+
HEMBA1004237	3	3.31	5.23	5.95	4.67	5.47		
HEMBA1004238	2.06	3.24	5.93	5.84	7.64	6.52		
HEMBA1004241	2.32	3.09	3.87	2.74	3.74	3.35		
HEMBA1004242	8.66	13.05	20.15	26.83	32.28	26.48	*	+
HEMBA1004243	1.8	2.09	3.58	2.8	3.03	2.76		
HEMBA1004246	1.6	2.68	5.65	6.18	6.24	6.15		
HEMBA1004247	0.89	2.73	3.74	3.69	4.23	3.37		
HEMBA1004248	4.01	3.54	3.85	5.91	8.31	7.47	**	+
HEMBA1004250	1.55	2.16	2.87	1.91	5.22	1.47		
HEMBA1004252	3.57	3.27	4.8	4.64	5.79	4.28		
HEMBA1004260	2.56	3.08	6.87	7.32	8.16	7.61		
HEMBA1004264	1.26	2.11	2.59	2.16	2.86	1.37		
HEMBA1004267	5.5	5.81	14.29	14.22	12.19	11.57		
HEMBA1004272	1.75	2.31	3.31	2.26	3.84	2.04		
HEMBA1004274	5.83	8.13	58.69	77.19	87.61	76.22	*	+
HEMBA1004275	1	5.4	3.34	1.49	4.49	2.42		
HEMBA1004276	2.27	2.2	3.42	3.45	4.2	3.06		
HEMBA1004279	2.13	2.33	4.37	3.29	5.2	3.88		
HEMBA1004284	1.78	2.56	6.03	4.16	4.9	5.23		
HEMBA1004286	1.41	1.35	2.44	1.65	3.55	2.1		
HEMBA1004289	2.58	4.17	5.59	5.16	7.18	8.44		
HEMBA1004293	20.24	18.64	51.03	77.3	52.39	74.25	*	+
HEMBA1004295	1.08	2.65	3.08	2.73	4.02	2.05		
HEMBA1004302	0.72	1.84	2.29	1.21	3.39	1.49		
HEMBA1004306	2.11	3.01	5.96	3.99	5.74	6.01		
HEMBA1004312	1.3	1.58	4.98	3.57	3.56	3.56		
HEMBA1004314	1.78	1.86	4.1	3.35	6.23	4.38		
HEMBA1004321	0.88	1.66	2.56	4.05	4.15	4.82	**	+
HEMBA1004323	2.4	3.16	4.7	4.77	5.29	5.1		
HEMBA1004327	1.18	1.78	3.23	3.51	4.46	3.1		
HEMBA1004329	5.57	6.73	15.22	16.29	15.33	15.14		
HEMBA1004330	3.93	3.54	4.06	5.87	7.54	5.36	*	+
HEMBA1004334	2.92	3.63	4.69	4.18	4.74	6.06		
HEMBA1004335	1.15	1.77	5.1	3.04	4.75	3.14		
HEMBA1004341	1.01	1.1	1.25	1.13	3.34	1.14		
HEMBA1004344	29.93	34.05	68.18	74.25	94.77	82.54	*	+
HEMBA1004347	0.67	1.65	2.21	2.31	2.25	2.78		
HEMBA1004349	12.07	11.55	22.26	19.39	22.71	24.41		
HEMBA1004352	2.06	2.56	6.96	6.05	6.92	6.58		
HEMBA1004353	10.21	14.95	25.3	19.64	26.63	25.83		

(1042)

特開2002-191363

2081

2082

HEMBA1004354	1.9	2.56	6.29	5.42	6.94	5.83		
HEMBA1004356	5.75	6.89	9.43	20.06	19.7	20.1	**	+
HEMBA1004360	1.35	1.16	2.73	1.23	4.23	2.26		
HEMBA1004366	1.97	1.91	3.57	4.53	6.9	4.73	*	+
HEMBA1004372	0.3	0.67	1.03	2.04	1.85	0.96		
HEMBA1004377	6.57	5.58	10.48	16.45	11.58	13.72	*	+
HEMBA1004389	8.39	7.87	13.69	16.87	19.3	11.15		
HEMBA1004391	1.18	0.88	1.37	1.27	4.23	2.34		
HEMBA1004393	65.85	84.9	114.17	77.61	53.19	85.8		
HEMBA1004394	0.84	1.14	1.88	1.68	3.9	2.59		
HEMBA1004396	1.62	0.82	3.24	2.43	5.44	2.15		
HEMBA1004401	4.33	4.37	5.25	8.95	14.42	11.05	*	+
HEMBA1004405	3.86	2.57	5.7	7.45	7.46	9.38	*	+
HEMBA1004408	4.27	2.66	5.34	7.19	8.8	9.85	*	+
HEMBA1004414	1.72	1.74	5.93	9.48	8.98	13.19	*	+
HEMBA1004429	2.95	3.96	4.23	4.68	6.9	6.8	*	+
HEMBA1004433	1.27	1.43	2.98	2.55	2.65	3.47		
HEMBA1004440	1.33	1.33	2.62	2.3	2.63	1.79		
HEMBA1004444	2.73	1.9	4.47	4.64	5.38	4.66		
HEMBA1004446	1.37	0.95	2.26	1.84	2.94	3.19		
HEMBA1004451	3.79	7.37	7.66	12.15	9.81	9.87	*	+
HEMBA1004452	0.71	1.96	2.89	3.53	8.91	2.36		
HEMBA1004454	1.56	1.71	3.06	2.81	4.83	4.1		
HEMBA1004460	1.91	3.49	7.49	4.39	6.91	5.19		
HEMBA1004461	1	1.87	1.43	1.09	3.95	1.99		
HEMBA1004468	3.22	4.71	7.36	9.92	12.73	8.79	*	+
HEMBA1004479	1.02	1.38	2.44	2.59	5.99	3.42		
HEMBA1004482	2.77	1.93	5.5	5.62	5.83	5.6		
HEMBA1004491	6.18	5.55	13	12.32	15.5	15.72		
HEMBA1004499	9.09	12.56	65.85	86.22	91.62	102.95	*	+
HEMBA1004502	1.81	1.9	4.82	3.14	4.83	2.49		
HEMBA1004505	1.57	2.42	4.6	4.78	7.66	4.09		
HEMBA1004506	0.96	2.32	3.48	3.28	5.58	2.73		
HEMBA1004507	29.33	30.94	74.36	85.26	112.08	83.25	*	+
HEMBA1004509	1.62	2.92	3.8	5.09	5.19	4.5	*	+
HEMBA1004523	1.04	1.68	1.65	1.82	3.64	3.02		
HEMBA1004528	9.41	10.46	59.94	86.06	89.35	98.27	*	+
HEMBA1004534	6.43	9.06	20.99	14.78	16.24	20.46		
HEMBA1004536	1.91	1.91	4.04	2.58	5.65	4.53		
HEMBA1004538	10.84	12.11	16.22	14.06	12.69	14.98		
HEMBA1004542	4.37	3.88	23.08	31.45	41.94	35.14	*	+
HEMBA1004552	2.98	1.35	4.27	9.89	9.44	7.1	**	+
HEMBA1004554	1.21	1.05	2.53	1.91	7.29	2.16		
HEMBA1004558	2.98	4.6	12.99	21.19	21.7	23.66	**	+
HEMBA1004560	1.74	2.06	4.16	3.47	6.18	4.25		
HEMBA1004564	4.31	3.3	8.73	7.27	9.96	8.65		
HEMBA1004566	50.51	49.26	83.43	126.94	55.59	138.13		
HEMBA1004573	1.07	1.97	3.91	2.91	4.11	2.3		
HEMBA1004576	3.68	3.22	10.49	4.73	5.64	3.11		
HEMBA1004577	6.11	7.2	14	11.74	13.04	10.71		

(1043)

特開 2002-191363

2083

2084

HEMBA1004586	1.56	1.32	5.09	5.01	6.65	4.67		
HEMBA1004596	3.97	3.19	19.1	27.4	39.04	32.16	*	+
HEMBA1004604	9.04	8.14	40.08	64.86	69.99	62.02	*	+
HEMBA1004607	0.96	1.97	4.17	3.02	4.39	3.96		
HEMBA1004610	1.16	1.89	3.19	2.91	4.8	2.64		
HEMBA1004617	0.78	0.89	3.29	2.77	4.53	2.51		
HEMBA1004622	1.05	2.39	5.1	4.07	5.84	5.04		
HEMBA1004626	1.56	2.36	4.75	4.06	4.93	2.33		
HEMBA1004629	1.5	1.34	3.95	3.17	4.58	2.74		
HEMBA1004631	1.57	1.73	4.41	3.3	5.73	4.97		
HEMBA1004632	1.02	1.3	3.43	2.38	3.64	2.36		
HEMBA1004633	3.2	3.98	8.84	9.3	9.04	10.74		
HEMBA1004636	1.29	2.07	3.22	3.23	3.61	6.58		
HEMBA1004637	1.57	2.12	4.19	3.97	4.85	2.39		
HEMBA1004638	1.31	1.67	3.26	1.8	4.24	1.52		
HEMBA1004645	3.04	2.88	6.5	5.07	6.21	5.08		
HEMBA1004656	4.38	2.76	4.96	4.13	4.58	3.39		
HEMBA1004657	16.78	17.12	35.48	31.85	17.55	22.74		
HEMBA1004666	1.27	2.2	3.32	2.78	4.52	1.44		
HEMBA1004669	2.49	3.6	6.16	7.32	8.34	5.55		
HEMBA1004670	3.1	2.74	6.27	5.11	6.96	3.3		
HEMBA1004672	1.29	2.33	4.85	2.58	7.25	3.63		
HEMBA1004689	23.54	21.34	82.29	90.21	98.37	106.84		
HEMBA1004690	4.74	5.24	15.26	19.89	22.39	20.76	*	+
HEMBA1004693	3.16	5.98	25.39	24.92	35.74	32.68		
HEMBA1004697	1.64	1.96	4.88	7.14	9.69	4.03		
HEMBA1004702	8.73	11.47	19.57	19.47	14.63	11.2		
HEMBA1004704	1.9	3.35	9.01	3.41	5.17	5.36		
HEMBA1004705	1.13	1.93	3.32	1.8	4.34	1.54		
HEMBA1004706	1.34	2.4	3.89	4.31	6.23	4.7		
HEMBA1004709	2.96	2.9	10.14	6.81	8.65	6.93		
HEMBA1004711	1.22	1.44	5.01	1.19	3.14	3.28		
HEMBA1004723	4.91	5.18	10.06	11.37	16.92	12.96	*	+
HEMBA1004725	4.14	4.9	7.68	5.67	8.62	8.54		
HEMBA1004730	3.57	2.83	4.74	3.64	8.59	4.41		
HEMBA1004733	1.07	2.64	2.62	1.06	3.54	2.45		
HEMBA1004734	2.57	3.22	4.46	2.68	5.28	5.32		
HEMBA1004736	1.1	1.89	7.12	3.87	12.1	5.09		
HEMBA1004748	2.24	0.94	5.6	3.02	5.22	2.83		
HEMBA1004749	6.88	8.33	19.68	16.49	19.03	23.29		
HEMBA1004751	1.96	1.76	5.55	3.99	9.86	5.04		
HEMBA1004752	1.51	1.6	4.23	4.56	4.11	3.32		
HEMBA1004753	29.15	25.19	85.53	59.89	62.5	95.58		
HEMBA1004755	7.02	6.32	12.37	9.73	12.72	14.63		
HEMBA1004756	1.45	1.76	3.86	2.34	5.21	2.17		
HEMBA1004758	1.18	1.64	4.53	3.92	5.65	3.17		
HEMBA1004763	1.79	2.39	5.56	5.45	6.53	6.09		
HEMBA1004768	0.83	1.64	2.89	1.69	4.26	1.38		
HEMBA1004770	1.09	1.36	2.43	1.47	3.53	1.94		
HEMBA1004771	0.99	1.02	2.44	2.18	3.26	2.57		

(1044)

特開 2002-191363

2085

2086

HEMBA1004775	4.07	3.84	7.29	8.61	9.62	13.74	*	+
HEMBA1004776	1.86	3.21	3.33	3.95	6.83	6.04	*	+
HEMBA1004778	1.75	2.24	6.11	3.21	7.64	5.74		
HEMBA1004784	1.51	1.59	3.11	2.36	4.18	3.62		
HEMBA1004785	1	1.78	4.15	2.18	6.54	4.48		
HEMBA1004789	2.34	2.07	4.42	1.87	5.64	2.87		
HEMBA1004795	0.62	1.89	3.13	2.45	4.23	1.42		
HEMBA1004797	1.06	0.84	1.85	2.31	2.85	2.76	*	+
HEMBA1004803	4.98	1.72	5.31	4.62	9.79	4.82		
HEMBA1004806	1.23	1.78	3.22	2.36	3.83	2.33		
HEMBA1004807	3.05	1.95	3.86	5.58	7.44	4.14		
HEMBA1004816	4.73	2.61	3.59	3.78	7.97	9.34		
HEMBA1004820	1.73	2.33	4.6	2.97	8.74	3.71		
HEMBA1004833	1.22	1.23	5.54	2.71	6.95	4.37		
HEMBA1004847	4.73	2.8	9.84	8	11.83	9.15		
HEMBA1004850	1.01	1.78	6.15	6.56	13.54	6.38		
HEMBA1004863	1.75	2.3	4.92	3.34	4.45	5.17		
HEMBA1004864	2.66	3.91	6.68	4.19	9.03	6.51		
HEMBA1004865	1.13	2.61	4.06	3.86	6.64	3.38		
HEMBA1004880	2.22	3.32	9.22	9.9	10.38	9.48		
HEMBA1004882	5.8	7.1	10.16	15.5	15.77	12.03	*	+
HEMBA1004885	2.34	6	8.67	8.58	8.67	9.27		
HEMBA1004889	3.25	3.24	6.5	7.67	10.74	9.8	*	+
HEMBA1004900	1.51	1.59	3.71	3.21	5.14	2.84		
HEMBA1004909	0.71	1.96	3.59	3.21	4.47	4.97		
HEMBA1004918	1.46	1.7	4.93	4.03	7.19	5.77		
HEMBA1004923	1.1	1.83	3.72	3.98	7.3	4.27		
HEMBA1004929	1.01	1.7	2.68	1.66	3.59	1.38		
HEMBA1004930	1.4	2.06	4.31	3.91	5.87	4.72		
HEMBA1004933	2.41	1.5	5.32	5.59	5.43	3.38		
HEMBA1004934	8.61	8.61	17.67	26.2	27.77	26.83	**	+
HEMBA1004937	1.36	2.42	3.57	2.79	7.51	2.54		
HEMBA1004943	1.11	1.75	3.63	2.56	4.88	3.68		
HEMBA1004944	0.95	2.51	4.73	3.85	4.79	3.58		
HEMBA1004946	4.78	4.38	12.5	12.69	14.08	12.8		
HEMBA1004952	0.9	1.83	4.04	2.76	4.41	2		
HEMBA1004954	3.55	3.21	7.92	6.02	6.4	6.12		
HEMBA1004956	1.37	1.11	2.78	2.12	3.66	1.81		
HEMBA1004960	0.86	0.75	3.27	4.17	6.72	3.06		
HEMBA1004971	2.41	2.49	6.81	6.45	8.05	6.92		
HEMBA1004972	2.57	2.31	3.63	2.07	5.66	5.84		
HEMBA1004973	1.16	1.78	3.46	2.53	3.27	3.18		
HEMBA1004977	3.04	3.08	5.36	5.34	6.29	6.36		
HEMBA1004978	4.53	5.08	14.93	21.43	19.01	25.37	*	+
HEMBA1004980	1.92	2.36	7.92	6.55	6.55	7.29		
HEMBA1004982	0.83	1.36	2.92	2.8	4.45	2.65		
HEMBA1004983	1.73	1.88	4.29	3.92	5.5	4.21		
HEMBA1004995	2.76	3.99	6.53	7.5	7.46	7.41		
HEMBA1005004	1.63	3.57	5.12	3.45	5.72	4.07		
HEMBA1005008	1.61	3.52	4.9	4.61	6.42	5.97		

(1045)

特開2002-191363

2087

2088

HEMBA1005009	4.55	3.9	8.37	11.95	6.96	9.22		
HEMBA1005019	3.1	2.57	5.78	6.6	8.07	7.19	*	+
HEMBA1005021	16.12	17.89	30.44	29.81	23.38	23.26		
HEMBA1005029	3.13	3.42	7.98	7.23	8.66	6.52		
HEMBA1005035	6.53	6.29	19.38	18.59	17.77	20.38		
HEMBA1005036	19.87	20.39	37.72	23.12	19.26	26.98		
HEMBA1005039	1.7	3.19	4.59	4.36	6.81	3.69		
HEMBA1005047	4.31	4.06	5.6	7.01	8.54	7.83	**	+
HEMBA1005050	2	2.93	5.07	4.33	6.76	4.39		
HEMBA1005062	2.48	3.06	5.62	2.85	5.13	2.03		
HEMBA1005066	2.28	2.55	5.72	4.81	8.94	6.89		
HEMBA1005067	5.81	10.37	17.44	25.77	30.77	26.84	*	+
HEMBA1005070	3.33	3.01	8.9	8.51	8.13	6.59		
HEMBA1005075	1.29	2.45	6.51	3.97	5.38	4.49		
HEMBA1005078	7.47	6.74	14.34	14.89	9.95	11.01		
HEMBA1005079	5.52	5.68	13.65	14.98	16.02	21.42		
HEMBA1005083	0.94	0.97	2.69	2.17	4.96	1.25		
HEMBA1005084	5.36	4.49	8.84	7.48	10.38	9.99		
HEMBA1005088	1.63	1.64	7.16	3.48	6.18	4.54		
HEMBA1005089	3.12	3.47	5.53	5.04	9	5.52		
HEMBA1005090	5.92	5.56	11.7	17.14	12.91	21.06	*	+
HEMBA1005096	0.88	2.47	3.98	3.35	3.94	2.88		
HEMBA1005101	2.29	2.08	4.54	3.23	5.6	4.54		
HEMBA1005107	1.2	1.92	3.2	2.25	4.48	2.3		
HEMBA1005113	0.96	2.18	3.35	2.12	5.53	2.72		
HEMBA1005123	3.35	3.46	10.83	9.05	10.41	8.28		
HEMBA1005133	2.6	2.26	7.59	5.11	4.58	7.09		
HEMBA1005135	1.19	2.77	3.35	1.47	5.28	3.69		
HEMBA1005145	5.84	6.38	12.56	13.06	14.71	17.84		
HEMBA1005149	4.21	3.2	7.92	6.51	8.54	8.47		
HEMBA1005152	1.81	3.06	3.59	3.31	5.05	3.39		
HEMBA1005159	1.76	1.96	3.62	2.2	8.12	2.2		
HEMBA1005172	120.6	113.48	240.76	174.65	194.57	222.24		
HEMBA1005185	3.16	2.05	4.47	3.54	6.7	4.15		
HEMBA1005186	1.55	2.65	3.96	3.52	8.39	2.03		
HEMBA1005195	2.15	1.08	3.76	1.93	6.73	2.24		
HEMBA1005201	3.27	3.45	8.02	5.52	7.17	8.91		
HEMBA1005202	5.93	4.58	9.4	8.29	9.76	12.83		
HEMBA1005204	86.99	67.86	141.56	134.68	133.57	172.39		
HEMBA1005206	6.39	5.08	66.26	72.89	92.07	89.66	*	+
HEMBA1005219	4.62	6.56	14.55	15.66	21.85	17.87	*	+
HEMBA1005223	2.4	1.91	4.92	1.7	3.83	3.86		
HEMBA1005229	1.18	2.32	3.12	1.57	4.88	2.32		
HEMBA1005230	1.22	1.56	4.24	2.71	5.92	4.32		
HEMBA1005232	0.86	1.13	1.6	0.91	3.81	2.17		
HEMBA1005238	0.84	1.77	2.67	2.16	3.27	2.29		
HEMBA1005241	2.77	2.68	7.42	7.63	11.38	7.62		
HEMBA1005244	0.87	1.97	5.63	2.57	8.33	6.19		
HEMBA1005246	8.84	7.84	17.94	12.96	18.21	14.79		
HEMBA1005251	3.06	2.57	6.81	6.6	8.47	7.24		

(1046)

特開 2002-191363

2089

2090

HEMBA1005252	2.3	2.7	5.31	4.73	11.4	5.01		
HEMBA1005267	2.77	3.31	7.25	6.77	12.66	9.89		
HEMBA1005274	1.75	1.53	3.28	4.8	6.38	6.51	**	+
HEMBA1005275	0.62	1.27	1.59	1.63	3.44	2.12		
HEMBA1005288	3.03	1.16	6.1	6.49	6.35	6.68		
HEMBA1005293	1.46	1.37	2.53	1.5	4.09	1.78		
HEMBA1005296	440.88	422.67	780.14	598.1	568.79	483.77		
HEMBA1005301	3.93	3.26	29.41	28.17	47.57	44.12		
HEMBA1005304	1.75	2.3	7	3.36	7.29	7.62		
HEMBA1005305	0.77	1.18	2.57	0.74	3.99	1.71		
HEMBA1005311	0.97	1.05	2.08	0.81	1.97	2.13		
HEMBA1005313	20.35	21.96	39.64	29.02	19.54	26.39		
HEMBA1005314	1	1.36	2.6	2.2	4.08	1.99		
HEMBA1005315	0.81	2.41	4.27	1.95	5.32	3.59		
HEMBA1005317	0.8	1.48	2.93	1.9	3.68	2.43		
HEMBA1005318	1.42	0.95	2.68	1.48	4.54	2.21		
HEMBA1005324	6.24	8.79	28.39	48.36	52.86	57.05	**	+
HEMBA1005331	2.77	5.64	15.05	21.73	18.68	21.33	*	+
HEMBA1005337	12.1	12.25	61.94	98.25	113.65	125.54	*	+
HEMBA1005338	4.53	4.54	27.15	34.04	35.53	42.4	*	+
HEMBA1005344	2.75	2.57	5.08	4.17	4.16	3.3		
HEMBA1005353	1.15	2.78	6.52	4.66	7.21	6.44		
HEMBA1005359	2.6	2.82	7.48	6.59	8.03	7.68		
HEMBA1005362	1.06	2.73	4.93	7.3	11.27	8.37	*	+
HEMBA1005364	0.96	1.48	3	2.55	3.78	2.75		
HEMBA1005367	0.96	1.53	4.19	3.3	4.21	2.99		
HEMBA1005372	1	2.24	4.18	2.3	5.1	3		
HEMBA1005374	2.29	3.31	7.83	6.4	6.45	8.09		
HEMBA1005379	3.19	3.61	7.65	29.57	25.57	24.46	**	+
HEMBA1005382	9.85	11.52	79.62	88.72	86.53	120.52		
HEMBA1005384	1.44	1.76	3.4	2.78	3.74	2.78		
HEMBA1005386	1.52	2.67	4.54	4.17	4.66	4.84		
HEMBA1005389	1.1	0.8	3.01	2.51	4.09	3.75		
HEMBA1005394	1.64	2.32	5	8.09	10.03	6.99	*	+
HEMBA1005403	4.19	4.24	8.81	8.46	11.37	8.87		
HEMBA1005408	1.43	1.71	5.3	4.52	7.12	6.96		
HEMBA1005410	1.06	1.74	3.22	1.78	4.97	2.42		
HEMBA1005411	1.38	2.58	4.54	2.84	4.09	2.74		
HEMBA1005423	4.14	5.87	4.72	12.75	8.59	11.04	*	+
HEMBA1005426	0.71	1.79	2.35	2.08	3.25	4.15		
HEMBA1005427	9.12	12.47	24.08	37.26	33.57	35.43	*	+
HEMBA1005430	1.71	1.43	3.23	3.26	4.07	4.78		
HEMBA1005438	1.42	2.33	4.26	4.01	5.41	2.66		
HEMBA1005443	15.57	16.88	35.83	20.61	27.42	29.93		
HEMBA1005447	1.55	1.95	3.42	2.25	5.9	3		
HEMBA1005449	1.5	1.36	3.29	1.82	5.83	3.21		
HEMBA1005452	8.96	10.79	63.65	107.56	94.42	105.84	*	+
HEMBA1005454	4.73	4.51	7.91	6.37	8.12	5.33		
HEMBA1005468	2.49	3.19	4	3.6	4.93	4.86		
HEMBA1005469	1.39	1.89	5.38	3.44	5.94	4.04		

(1047)

特開 2002-191363

2091

2092

HEMBA1005472	4.04	4.76	7.81	6.79	6.75	8.2		
HEMBA1005474	4.91	5.42	11.67	7.77	9.4	9.23		
HEMBA1005475	7.8	7.09	11.75	15.34	10.75	13.59		
HEMBA1005489	2.47	3.48	4.94	6.34	7	6.54	*	+
HEMBA1005497	0.65	1.97	3.47	2.86	4.42	2.14		
HEMBA1005500	4.64	4.25	11.48	9.17	13.98	10.59		
HEMBA1005506	3.88	2.15	4.02	2.71	3.57	1.29		
HEMBA1005508	5.79	7.67	13.3	11.65	14.52	10.99		
HEMBA1005511	2.97	2.29	9	4.36	6.36	6.14		
HEMBA1005513	6.3	9.05	55.46	50.46	60.73	57.45		
HEMBA1005517	1.8	2.89	3.88	4.37	6.41	3.74		
HEMBA1005518	1.48	2.49	3.94	2.94	5.22	2.3		
HEMBA1005520	1.89	3.19	8.8	8.37	10.13	9.32		
HEMBA1005522	1.78	2.29	3.37	2.52	4.72	1.45		
HEMBA1005526	3.24	2.8	6.25	4.37	7.26	4.58		
HEMBA1005528	8.59	16.13	16.28	20.24	22.18	20.01		
HEMBA1005530	2.28	3.62	5.13	4.91	7.75	4.51		
HEMBA1005538	11.07	10.7	15.34	8.46	4.44	5.89	*	-
HEMBA1005539	30.73	31.96	63.32	50.52	49.29	34.26		
HEMBA1005545	1.2	1.08	3	4.28	6.22	2.25		
HEMBA1005548	2.38	2.88	11.1	11.04	12.72	9.21		
HEMBA1005552	2.76	4.58	12.05	9.13	12.8	10.25		
HEMBA1005558	1.59	2	5.86	2.18	6.97	4.17		
HEMBA1005568	3.11	2.96	8.24	6.28	8.45	7.5		
HEMBA1005570	1.87	2.75	3	1.96	4.49	7.13		
HEMBA1005576	1.74	2.8	2.4	3.03	3.55	3.57	*	+
HEMBA1005577	1.14	2.02	2.78	0.95	3.08	0.87		
HEMBA1005581	4.29	4.22	6.85	14.27	18.37	8.15		
HEMBA1005582	2.74	2.57	5.35	3.25	7.91	3.21		
HEMBA1005583	2.47	3.31	5.86	3.32	6.43	3.44		
HEMBA1005588	2.51	2.85	6.27	5	6.54	6.14		
HEMBA1005593	1.5	1.4	2.85	1.83	4.57	2.89		
HEMBA1005595	2.62	2.82	4.15	3.31	4.65	4.76		
HEMBA1005597	4.77	5.18	8.13	9.39	10.34	8.64	*	+
HEMBA1005606	2.29	2.76	5.79	3.96	6.91	5.83		
HEMBA1005609	2.84	2.64	6.61	4.19	5.77	6.81		
HEMBA1005616	2.01	1.66	8.03	5.44	8.75	7.03		
HEMBA1005621	2.43	1.91	4.42	4.13	6.24	2.7		
HEMBA1005627	3.84	3.92	11.61	9.73	15.14	14.89		
HEMBA1005628	12.1	12.91	20.55	17.92	23.35	18.08		
HEMBA1005631	13.47	11.94	26.82	22.77	22.87	29.03		
HEMBA1005632	1.33	2.33	5.06	3.14	3.68	4.47		
HEMBA1005634	3.06	3.42	5.15	2.81	7.68	5.18		
HEMBA1005662	1.18	1.27	3.17	1.06	4.57	3.03		
HEMBA1005666	5.89	4.51	10.09	10.5	9.01	10.25		
HEMBA1005670	1	1.08	4.06	2.87	4.35	3.19		
HEMBA1005671	2.11	3.38	5.07	5.36	9	5.9		
HEMBA1005679	2.33	4.64	7.39	6.5	10.44	10.19		
HEMBA1005680	2.63	2.14	5.9	5.51	7.59	7.72		
HEMBA1005685	2	1.89	7.27	3.8	6.73	1.97		

(1048)

特開 2002-191363

2093

2094

HEMBA1005698	5.96	4.75	12.88	11.78	14.17	9.93		
HEMBA1005699	1.4	1	2.45	2.17	3.66	2.96		
HEMBA1005703	1.22	1.27	3.57	1.79	3.56	1.88		
HEMBA1005705	2.39	2.78	6.45	3.41	6.27	3.89		
HEMBA1005712	1.23	1.34	4.52	2.18	4.84	2.37		
HEMBA1005717	1.55	1.89	4.7	1.34	5.36	2.16		
HEMBA1005718	5.27	4.35	7.8	10.09	10	15.72	*	+
HEMBA1005721	15.93	20.34	26.12	37.74	25.37	32.79		
HEMBA1005722	18	19.32	35.72	25.36	30.28	26.91		
HEMBA1005724	2.17	2.05	4.6	2.47	6.56	2.98		
HEMBA1005732	1.33	1.54	4.89	7.22	8.38	5.67	*	+
HEMBA1005737	1.49	1.19	2.95	2.44	4.57	1.75		
HEMBA1005742	3.4	4.65	5.7	4.43	6.35	4.22		
HEMBA1005746	1.2	1.61	3.23	5.69	4.42	4.85	*	+
HEMBA1005747	3.8	3.51	7.52	6.08	6.55	6.67		
HEMBA1005749	9.64	9.37	17.99	21.62	19.82	18.15		
HEMBA1005755	0.49	2.5	3.35	1.66	4.41	2.9		
HEMBA1005760	1.59	2.1	5.37	3.7	4.86	4.14		
HEMBA1005765	1.47	2.89	5.97	5.91	7.42	7.21		
HEMBA1005766	8.52	10.49	61.64	93.4	112.09	106.28	*	+
HEMBA1005780	4.98	5.15	14.49	13.93	16.26	12		
HEMBA1005795	1.15	1.8	3.71	4.28	5.45	4.26	*	+
HEMBA1005809	9.13	9.7	20.95	14.79	29.83	22.75		
HEMBA1005813	1.76	2.74	6.74	5.39	9	7.08		
HEMBA1005815	0.62	1.86	2.75	2.74	4.78	2.13		
HEMBA1005822	1.34	2.83	5.82	3.77	6.65	5.12		
HEMBA1005829	1.49	2.66	4.94	5.11	4.53	5.52		
HEMBA1005833	1.83	1.28	4.41	3.89	5.17	4.57		
HEMBA1005834	2.53	1.87	4.84	5.48	11.84	6.81		
HEMBA1005844	27.89	31.06	52.5	62.42	59.49	58.73	*	+
HEMBA1005852	5.75	4.18	10.12	10.15	12.34	11.51		
HEMBA1005853	2.81	3.05	9.32	5.68	13.84	9.26		
HEMBA1005878	4.42	4.52	10.14	8.84	10.05	11.01		
HEMBA1005883	1.67	2.81	4.56	2.81	5.43	4.01		
HEMBA1005884	1.87	2	3.21	2.27	5.9	2.28		
HEMBA1005891	2.39	3.04	3.59	4.34	5.46	3.95		
HEMBA1005894	2.12	2.41	8.68	8.23	9.08	6.9		
HEMBA1005898	4.52	3.75	7.4	10.51	10.74	9.53	*	+
HEMBA1005902	2.57	3.14	8.63	5.36	6.75	5.82		
HEMBA1005907	1.15	2.28	2.2	2.28	4.2	1.74		
HEMBA1005909	0.93	2.68	2.97	2.96	4.55	2.6		
HEMBA1005911	1.66	3.12	4.9	3.96	9.5	3.41		
HEMBA1005912	8.83	7.86	13.57	16.44	8.16	15.58		
HEMBA1005913	5.05	5.39	10.39	9.57	7.71	7.06		
HEMBA1005921	2.36	3.86	6.29	5.25	10.06	7.64		
HEMBA1005922	5.49	5.47	6.99	8.51	10.1	8.74	*	+
HEMBA1005929	1.91	2.53	7.53	5.05	5.68	6.47		
HEMBA1005931	3.32	2.95	6.04	5.31	9.49	6.09		
HEMBA1005934	2.9	4.69	8.2	8.17	10.14	9.73		
HEMBA1005945	3.1	4.12	7.73	6.36	9.39	6.65		

(1049)

特開 2002-191363

2095

2096

HEMBA1005962	1.72	1.81	3.16	1.93	3.96	1.72		
HEMBA1005963	1.86	1.67	3.91	2.26	3.95	2.05		
HEMBA1005990	6.04	7.39	17.86	14.14	16.18	17.02		
HEMBA1005991	2.39	3.35	8.55	7.2	7.64	6.94		
HEMBA1005999	2.34	4.39	8.84	6.52	8.18	8.75		
HEMBA1006002	3.53	5	7.73	5.96	10.24	8.99		
HEMBA1006005	0.96	2.31	3.93	2.87	5.65	3.27		
HEMBA1006011	26.27	24.03	34.08	62.88	44.8	64.92	*	+
HEMBA1006013	2.43	2.57	6.31	4.17	6.87	3.74		
HEMBA1006016	1.65	1.98	5.64	3.22	5.34	2.16		
HEMBA1006019	2.97	3.23	6.91	4.46	7.22	5.44		
HEMBA1006021	5.06	6.45	9.21	10.29	12.48	8.77		
HEMBA1006022	3.19	4.34	6.89	6.52	6.1	5.94		
HEMBA1006031	1.32	2.46	4.38	3.72	5.57	5.34		
HEMBA1006035	3.05	3.72	7.96	4.99	6.37	4.54		
HEMBA1006036	2.02	2.3	7	4.27	10.14	7.22		
HEMBA1006042	3.36	3.1	8.51	5.76	9.59	8.31		
HEMBA1006044	1.44	1.99	3.61	2.8	3.4	1.67		
HEMBA1006045	1.98	1.99	5.08	4.3	8.55	5.64		
HEMBA1006048	2.42	4.18	5.41	6.84	7.96	8.18	*	+
HEMBA1006053	1.51	2.72	3.55	3.09	3.3	3.93		
HEMBA1006055	1.84	1.91	2.46	3.04	4.92	3.67	*	+
HEMBA1006058	4.04	4.62	11.59	7.42	10.66	12.41		
HEMBA1006063	9.2	9.36	32.62	26.37	33.39	27.19		
HEMBA1006067	4.14	3.27	5.81	5.8	8.4	7.17		
HEMBA1006081	0.84	2.59	4.77	2.54	7.08	2.49		
HEMBA1006089	2.58	4.48	6.82	8.28	10.12	9.66	*	+
HEMBA1006090	1.66	2.31	2.28	1.66	5.26	1.84		
HEMBA1006091	1.1	1.35	1.75	3.15	4.31	2.95	*	+
HEMBA1006093	1.65	1.77	4.21	2.27	6.67	4.26		
HEMBA1006099	11.9	9.88	23.12	21.57	16.43	20.57		
HEMBA1006100	2.78	3.18	13.25	8.38	15.71	13.52		
HEMBA1006108	2.69	2.08	4.22	3.42	5.24	3.75		
HEMBA1006114	7.21	8.76	39.36	34.51	62.97	50.98		
HEMBA1006121	1.18	1.8	3.13	2.43	5.83	2.91		
HEMBA1006124	1.79	1.74	4.47	3.11	4.89	4.13		
HEMBA1006125	18.52	14.19	23.44	25.23	22.56	34.45		
HEMBA1006130	5.15	3.1	7.57	6.89	7.84	9.51		
HEMBA1006138	2.43	2.41	5.55	5.27	9.2	8.12		
HEMBA1006142	2.62	1.87	6.23	6.67	9.36	8.91	*	+
HEMBA1006150	8.32	7.44	21.06	20.42	21.82	24.68		
HEMBA1006151	567.67	524.4	796.07	915.23	875.11	682.15		
HEMBA1006155	0.93	1.33	2.92	1.44	5.54	1.94		
HEMBA1006158	3.06	4.95	7.5	5.7	9.04	6.31		
HEMBA1006164	2.61	1.96	5.89	5.26	4.54	5.82		
HEMBA1006171	29.76	24.08	54.44	32.3	33.25	34		
HEMBA1006173	5.15	3.15	30.41	36.16	55.46	57.62	*	+
HEMBA1006176	315.12	232.27	427.53	476.77	458.01	381.49		
HEMBA1006182	1.47	1.71	6.13	3.02	5.84	5.82		
HEMBA1006197	6.14	5.09	9.26	9.53	12.1	10.41		

(1050)

特開2002-191363

2097

2098

HEMBA1006198	10.07	6.46	26.71	35.03	45.64	55.68	*	+
HEMBA1006213	1.98	1.78	2.4	2.33	2.69	2.3		
HEMBA1006217	44.9	41.62	72.62	94.44	74.92	78.25		
HEMBA1006226	40.86	36.82	63.27	77.37	79.96	57.39		
HEMBA1006235	2.13	1.96	4.34	2.6	5.85	2.32		
HEMBA1006248	1.74	2.03	4.04	3.37	6.4	2.95		
HEMBA1006251	5.41	7.29	7.88	11.96	12.95	9.22	*	+
HEMBA1006252	0.72	1.28	3.67	2.25	3.56	2.92		
HEMBA1006253	2.13	2.26	4.1	5.93	5.38	5.76	*	+
HEMBA1006259	1.96	2.29	6.02	5.7	3.9	5.32		
HEMBA1006261	12.25	7.27	18.84	14.23	14.08	13.61		
HEMBA1006268	2.05	2.56	3.86	4.83	5.09	4.06	*	+
HEMBA1006271	2.04	3.99	10.58	7.13	9.33	8.51		
HEMBA1006272	0.97	2.26	2.84	2.38	6.01	1.93		
HEMBA1006273	1.53	2.09	4.55	3.58	4.52	2.46		
HEMBA1006276	2.8	1.26	3.62	4.45	5.84	2.82		
HEMBA1006278	1.57	2.03	3.19	4.08	4.51	2.72		
HEMBA1006283	3.09	3	6.08	7.34	11.13	7.09		
HEMBA1006284	2.47	1.57	3.14	4.75	6.96	2.82		
HEMBA1006291	1.42	2.56	4.41	4.6	6.16	2.57		
HEMBA1006292	3.36	5.12	17.34	19.95	23.83	21.11	*	+
HEMBA1006293	1.83	1.46	3.19	2.92	4.02	1.36		
HEMBA1006299	1.92	2.26	7.03	5.02	6.39	4.74		
HEMBA1006309	2.26	1.43	3.53	4.47	4.69	3.73	*	+
HEMBA1006310	4.14	4.32	7.72	9.51	8.34	5.7		
HEMBA1006311	1.4	2.33	6.68	5.59	5.85	5.72		
HEMBA1006313	1.2	1.6	2.74	4.29	4.68	2.26		
HEMBA1006316	2.16	3.08	6.72	6.63	6.28	5.59		
HEMBA1006328	2.78	4.28	13.48	15.14	17	16.56		
HEMBA1006334	1.46	3.1	2.77	2.79	3.95	2.05		
HEMBA1006335	10.42	13.98	21.66	20.81	18.03	16.77		
HEMBA1006344	2.86	3.55	6.9	7.18	6.88	5.75		
HEMBA1006347	2.04	1.83	4.2	2.94	2.76	1.68		
HEMBA1006349	2.47	2.79	6.73	3.62	5.65	3.5		
HEMBA1006352	1.65	1.65	3.27	2.63	5.64	1.32		
HEMBA1006357	4.99	4.26	8.6	8.36	7.8	7.74		
HEMBA1006358	1.67	2.57	4.95	4.7	5.11	5.63		
HEMBA1006359	1.56	2.17	4.02	3.67	5.48	3.3		
HEMBA1006360	2.53	2.12	5.1	9.73	10.02	8.61	**	+
HEMBA1006364	1.71	2.76	4.82	4.02	5.55	3.51		
HEMBA1006377	5.8	8.03	13.25	20.48	16.19	15.31	*	+
HEMBA1006380	1.31	1.57	8.75	6.78	7.01	6.9		
HEMBA1006381	2.38	3.07	12.65	6.18	7.97	6.66		
HEMBA1006385	3.21	3.33	8.65	5.41	8.17	5.32		
HEMBA1006390	9.49	7.85	14.66	22.01	20.52	22.7	**	+
HEMBA1006391	6.58	6.85	6.73	12.83	10.12	13.15	**	+
HEMBA1006398	1.32	1.67	4.19	2.57	4.12	1.69		
HEMBA1006405	23.81	23.5	38.82	23.85	22.13	24.92		
HEMBA1006410	8.26	4.16	6.14	5.74	11.61	6.06		
HEMBA1006416	2.14	2.62	5.93	5.3	6.98	4.23		

(1051)

特開2002-191363

2099

2100

HEMBA1006418	5.06	5.49	11.76	8.17	8.09	5.92		
HEMBA1006419	2.67	3.93	8	6.89	7.77	5.2		
HEMBA1006421	2.03	3.28	3.09	3.44	4.19	2.27		
HEMBA1006424	1.48	1.92	3.59	1.94	5.42	1.84		
HEMBA1006426	3.03	3.99	7.91	7.23	7.87	5.51		
HEMBA1006430	2.31	2.64	6.29	5.89	7.43	5.08		
HEMBA1006438	2.06	2.22	6.35	4.37	6.53	2.92		
HEMBA1006445	1.98	2.68	5.72	6.11	5.6	4.16		
HEMBA1006446	1.32	2.61	5.59	2	6.51	2.43		
HEMBA1006456	3.51	5.07	8.64	14.9	21.76	15.91	*	+
HEMBA1006461	1.54	2.18	5.35	4.35	5.49	4.07		
HEMBA1006467	1.52	1.78	3.61	2.82	7.24	3.2		
HEMBA1006470	4.06	4.03	22.46	18.72	29.54	19.52		
HEMBA1006471	1.58	1.6	6.6	6.45	6.78	6.26		
HEMBA1006474	7.35	6.37	43.12	55.87	62.01	52.31	*	+
HEMBA1006476	9.48	10.05	66.66	94.3	119.21	92.71	*	+
HEMBA1006482	71.42	71.44	219.31	199.96	180.73	192.88		
HEMBA1006483	2.03	2.96	7.22	3.36	4.88	3.24		
HEMBA1006485	2.24	1.72	6.31	4.09	6.53	5.51		
HEMBA1006486	5.08	5.55	13.73	18.4	16	15.46	*	+
HEMBA1006489	1.21	2.18	4.1	3.18	5.4	2.17		
HEMBA1006492	5.34	7.71	10.69	15.16	24.33	15.92	*	+
HEMBA1006494	1.27	1.18	2.67	2.44	3.88	1.86		
HEMBA1006497	1.67	2.33	4.75	4.3	3.79	4.49		
HEMBA1006501	7.61	7.52	62.05	58.03	78.37	58.59		
HEMBA1006502	4.73	3.55	15.72	18.66	22.55	21.66	*	+
HEMBA1006507	8.7	6.8	51	49.69	71.2	48.41		
HEMBA1006517	1.51	1.99	5.07	3.43	5.9	4.64		
HEMBA1006521	1.79	1.8	4	2.41	4.55	3.02		
HEMBA1006529	4.77	3.74	3.86	5.97	5.83	2.78		
HEMBA1006530	1.8	1.39	2.06	1.62	3.53	2.42		
HEMBA1006535	1.66	1.43	2.01	2.66	2.81	2.43	**	+
HEMBA1006536	0.59	2.22	3.96	3.04	3.23	2.33		
HEMBA1006540	1.61	1.68	3.33	3.05	4.1	3.56		
HEMBA1006544	1.39	1.63	8	3.54	5.85	4.35		
HEMBA1006546	2.06	2.56	6.98	4.25	5.77	4.51		
HEMBA1006549	1.74	2.13	5.93	4.57	4.63	4.61		
HEMBA1006559	2.55	1.45	4.63	2.99	5.76	3.32		
HEMBA1006562	0.74	1.32	4.07	2.39	5.24	2.72		
HEMBA1006566	0.67	1.28	0.97	1.34	1.69	0.99		
HEMBA1006569	2.33	1.36	3.97	3.25	3.89	4.02		
HEMBA1006572	1.02	2.38	2.94	2.68	3.92	2.01		
HEMBA1006579	20.44	16.82	51.93	59.14	66.28	72.54	*	+
HEMBA1006583	3.17	2.85	5.59	5.83	5.94	6.45		
HEMBA1006595	1.82	1.63	4.29	2.83	5.22	3.68		
HEMBA1006597	1.65	2.1	5.34	3.81	6.75	3.77		
HEMBA1006606	1.75	2.31	4.96	3.72	5.71	2.8		
HEMBA1006612	2.63	3.54	7.11	7.32	5.31	7.07		
HEMBA1006617	1.93	2.58	6.24	4.62	4.6	5.41		
HEMBA1006624	6.37	8.61	16.93	12.84	15.78	10.91		

(1052)

特開2002-191363

2101

2102

HEMBA1006631	3.24	3.13	8.01	8.11	11.54	6.97		
HEMBA1006635	1.7	2.57	5.7	4.27	6.27	3.8		
HEMBA1006639	1.12	1.98	4.52	4.14	5.3	3.06		
HEMBA1006643	1.88	1.23	1.85	2.12	3.81	2.17		
HEMBA1006648	7.69	6.84	16.92	17.63	23.57	23.73	*	+
HEMBA1006652	5.96	7.86	16.11	16.41	16.83	17.34		
HEMBA1006653	2.21	3.66	7.23	6.92	5.18	5.17		
HEMBA1006658	5.04	6.58	12.68	14.07	13.59	12.01		
HEMBA1006659	6.8	10.14	48.92	66.47	78.17	67.44	*	+
HEMBA1006665	1.44	0.89	3.32	2.77	2.91	1.82		
HEMBA1006666	1.83	1.25	3.23	3.24	4.14	2.45		
HEMBA1006671	8.69	6.74	11.66	17.25	16.13	15.21	**	+
HEMBA1006674	1.64	1.99	7.42	6.71	9.73	7.34		
HEMBA1006676	1.46	2.36	5.19	4.28	6.75	3.19		
HEMBA1006682	2.17	1.64	3.43	2.2	4.98	1.88		
HEMBA1006688	1.48	2.46	4.74	3.31	4.38	3.54		
HEMBA1006695	1.58	2.41	4.85	3.54	5.79	2.46		
HEMBA1006696	2.84	3.93	6.29	5.95	6.74	5.82		
HEMBA1006702	3.31	1.83	13.28	4.13	5.58	4.87		
HEMBA1006707	2.89	2.62	5.9	5.04	7.54	5.5		
HEMBA1006708	2.21	1.52	5.71	4.42	4.57	2.63		
HEMBA1006709	1.64	1.97	4.6	4.29	5	4.74		
HEMBA1006717	1.58	2.28	3.58	2.5	4.93	2.28		
HEMBA1006724	2.68	3.42	4.55	4.45	4.5	5.47		
HEMBA1006731	1.83	2.95	3.95	4.12	5.51	3.1		
HEMBA1006737	1.82	3.5	6.59	3.89	5.09	4.45		
HEMBA1006742	1.78	2.44	4.16	3.32	4.14	3.61		
HEMBA1006743	4	4.02	11.48	14.16	17.25	11.88		
HEMBA1006744	1.84	1.79	7.74	6.6	8.29	5.72		
HEMBA1006749	1.14	1.27	3.72	1.88	3.8	1.71		
HEMBA1006752	16.53	16.28	26.81	35.31	18.85	33.99		
HEMBA1006754	1.44	2.6	3.63	5.55	4.33	2.49		
HEMBA1006758	1.38	2.83	4.25	6.89	4.68	4.02		
HEMBA1006767	3	4.14	7.88	6.81	7.89	5.51		
HEMBA1006770	5.05	2.61	7.12	7.89	8.1	5.05		
HEMBA1006779	4.44	4.1	10.99	9.57	10.28	8.42		
HEMBA1006780	3.28	3.19	10.27	7.6	8.33	8.38		
HEMBA1006789	2.83	1.87	11.34	4.55	5.42	5.51		
HEMBA1006795	2.13	2.45	7.58	4.56	7.87	3.29		
HEMBA1006796	4.31	3.15	5.37	5.85	7.06	4.72		
HEMBA1006805	2.72	2.73	5.9	12.77	17.38	12.96	**	+
HEMBA1006807	30.32	28.07	75.38	54.22	54.72	67.82		
HEMBA1006813	0.93	1.73	2.81	1.93	4.17	0.97		
HEMBA1006819	3.73	4.53	8.5	7.02	8.09	5.29		
HEMBA1006821	1.56	2.37	6.09	5.05	4.24	4.36		
HEMBA1006824	2.13	3.13	7.39	5.69	6.04	5.75		
HEMBA1006832	19.84	18.63	56.97	59.26	61.56	53.28		
HEMBA1006834	13.23	12.47	20.38	29.88	28.37	21.62	*	+
HEMBA1006835	1.11	1.49	3.88	5.08	7.25	4.6	*	+
HEMBA1006843	19.27	17.89	35.47	55.34	39.67	68.17	*	+

(1053)

特開 2002-191363

2103

2104

HEMBA1006849	5.64	4.37	11.23	10.07	11.23	10.83		
HEMBA1006850	31.45	33.76	60.2	44.62	53.25	45.59		
HEMBA1006861	12.19	11.3	24.61	22.49	16.43	17.99		
HEMBA1006865	5.42	8.35	31	33.77	33.36	34.49		
HEMBA1006867	4.32	5.03	6.41	6.27	7.76	6.15		
HEMBA1006873	3.14	3.59	8.87	6.5	9.26	7.75		
HEMBA1006877	2.52	4.03	5.87	4.03	6.26	3.68		
HEMBA1006878	2.52	2.36	6.79	3.82	7.86	3.51		
HEMBA1006879	6.19	5.68	8.83	11.06	15.86	17.33	*	+
HEMBA1006884	10.11	3.17	6.59	6.15	8.42	8.78		
HEMBA1006885	7.02	6.82	14.16	20.86	19.11	21.73	*	+
HEMBA1006886	20.38	17.25	26.55	29.45	29.15	40.93		
HEMBA1006889	2.61	4.02	4.85	4.48	5.99	6.09		
HEMBA1006896	19.11	24.76	31.7	33.67	39.11	40.41	*	+
HEMBA1006900	6.19	5.93	20.76	16.81	23.73	18.45		
HEMBA1006902	1.43	2.45	3.86	4.03	6	3.98		
HEMBA1006912	1.24	1.74	6.86	4.12	5.8	5.3		
HEMBA1006914	6.64	6.11	18.27	14.81	18.62	15.03		
HEMBA1006916	3.11	2.71	5.78	10.29	7.48	9.36	*	+
HEMBA1006921	3.03	3.5	9.63	9.77	11.26	13.59		
HEMBA1006926	2.65	2.61	5.68	5.01	6.53	6.98		
HEMBA1006927	3.06	2.2	5.17	3.57	5.26	5.89		
HEMBA1006929	2.94	2.69	4.02	4.31	6.36	5.25	*	+
HEMBA1006936	3.72	3.21	6.51	4.67	6.25	5.45		
HEMBA1006938	1.21	2.11	6.57	2.37	3.76	3.44		
HEMBA1006941	9.52	8.15	12	19.26	28.62	23.74	**	+
HEMBA1006942	5.2	2.63	6.65	10.7	10.65	11.4	**	+
HEMBA1006945	10.07	5.91	16.81	23.73	17.09	19.91		
HEMBA1006949	1.6	1.43	3.88	2.48	5.34	2.81		
HEMBA1006952	1.16	1.66	2.98	3.02	5.04	2.22		
HEMBA1006960	2.53	2.78	7.66	5.9	8.28	8.68		
HEMBA1006973	1.74	2.27	5.91	4.7	7.84	5.54		
HEMBA1006974	2.49	3.44	6.76	6.09	11.01	8.14		
HEMBA1006976	1.39	1.5	4.12	3.18	4.96	4.36		
HEMBA1006989	1.85	1.66	6.51	2.05	3.01	1.81		
HEMBA1006993	2.71	2.39	6.49	6.11	7.69	8.79		
HEMBA1006996	0.74	1.15	2.98	2.52	3.13	3.58		
HEMBA1007001	1.91	2.47	5.12	3.98	6.37	4.76		
HEMBA1007002	7.02	4.12	31.4	26.92	38.45	42.11		
HEMBA1007013	1.02	0.94	3.04	1.44	4.39	2.27		
HEMBA1007016	2.02	1.43	5.06	3.27	5.97	5.28		
HEMBA1007017	0.69	1.24	2.55	1.42	3.33	1.83		
HEMBA1007018	4.02	4.52	6.54	7.65	6.1	7.07		
HEMBA1007044	8.13	8.41	17.4	15.48	11.97	12.27		
HEMBA1007045	1.64	2.15	4.42	2.61	5.08	3.47		
HEMBA1007051	2.26	2.56	4.71	3.42	4.42	3.28		
HEMBA1007052	2.23	1.25	3.47	2.37	4.62	1.83		
HEMBA1007053	1.83	3.14	4.03	2.64	4	2.5		
HEMBA1007057	0.92	2.56	3.21	3.52	4.03	3.41		
HEMBA1007062	0.91	0.82	2.73	2.34	2.87	1.82		

(1054)

特開2002-191363

2105

2106

HEMBA1007063	3.87	2.7	8.87	8.56	8.02	7.58		
HEMBA1007066	1.72	2.03	3.62	2.2	4.39	2.71		
HEMBA1007069	1.36	2.29	3.87	3.84	3.48	3.66		
HEMBA1007073	1.93	2.16	6.12	2.96	9.57	3.6		
HEMBA1007076	1.48	2.6	5.56	4.66	7.6	3.4		
HEMBA1007078	6.5	6.83	18.61	26.96	27.47	23.78	*	+
HEMBA1007080	7.6	9.46	38.27	61.02	68.15	51.34	*	+
HEMBA1007084	1.28	1.42	4.76	3.68	6.27	4.76		
HEMBA1007085	3.28	2.89	8.21	5.76	7.21	5.42		
HEMBA1007087	2.6	2.88	6.31	3.96	6.92	5.63		
HEMBA1007089	26.17	28.1	43.8	46.11	34.41	29.42		
HEMBA1007095	75.81	62.79	111.43	134.53	48.47	121.4		
HEMBA1007101	2.78	3.27	8	26.73	21.52	19.57	**	+
HEMBA1007104	1.87	1.92	3.52	2.46	4.53	2.57		
HEMBA1007106	4.77	4.8	9.03	16.42	12.5	9.49		
HEMBA1007112	3.01	3.07	5.16	6.39	6	4.77		
HEMBA1007113	1.53	2.29	9.04	6.03	6.1	5.97		
HEMBA1007121	13.76	14	92.08	116.14	111.53	129.12	*	+
HEMBA1007129	1.54	2.44	2.87	2.66	4.37	1.89		
HEMBA1007147	1.68	2.88	4.4	3.96	4.76	4.06		
HEMBA1007149	5.3	7.24	8.38	10.48	6.82	9.73		
HEMBA1007151	0.85	1.87	3.38	3.32	3.88	2.54		
HEMBA1007172	1.26	1.91	4.13	2.96	4.81	3.51		
HEMBA1007174	1.4	1.43	2.75	3.96	3.65	2.5		
HEMBA1007176	2.58	3.95	11.7	6.7	6.78	4.52		
HEMBA1007178	4.77	4.71	9.32	10.94	13.03	8.12		
HEMBA1007185	9.38	10.32	9.59	19.5	7.83	15.16		
HEMBA1007186	1.71	2.76	4.49	4.95	5.47	3.86		
HEMBA1007194	4.81	3.43	5.58	7.83	9.34	8.67	**	+
HEMBA1007200	1.18	2.33	2.9	3.25	4.6	1.66		
HEMBA1007203	1.54	3.5	5.38	6.03	6.87	5.05		
HEMBA1007206	1.92	2.46	5.72	7.07	7.91	5.94		
HEMBA1007224	5.4	6.5	9.06	9.23	5.85	8.06		
HEMBA1007226	7.19	8.07	40.61	59.31	70.51	62.19	*	+
HEMBA1007240	10	10.96	13.45	15.35	7.71	11		
HEMBA1007241	3.59	2.88	4.56	4.61	6.81	3.63		
HEMBA1007242	2.52	2.86	5.01	6.29	6.87	4.23		
HEMBA1007243	10.23	10.91	69.57	70.17	95.69	82.75		
HEMBA1007251	1.32	1.8	4.14	3.02	3.67	2.01		
HEMBA1007256	1.39	1.91	3.36	3.93	5.74	3.44		
HEMBA1007267	3.19	3.71	8.75	8.73	9.18	8.1		
HEMBA1007273	0.98	2.66	3.84	3.56	5.82	2.25		
HEMBA1007279	1.55	2.25	3.52	2.95	4.35	2.02		
HEMBA1007281	1.73	1.54	2.12	2.95	4.43	1.01		
HEMBA1007283	2.45	3.15	6.78	6.37	6.58	5.96		
HEMBA1007288	2.12	2.77	5.54	4.35	6.74	5.48		
HEMBA1007291	1.59	1.8	4.29	2.14	4.4	0.98		
HEMBA1007299	20.39	22.25	39.67	40.95	47.97	40.26		
HEMBA1007300	2.08	2.75	3.59	4.17	4.45	4.07	*	+
HEMBA1007301	1.97	2.82	3.15	3.73	3.99	3.44	*	+

(1055)

特開2002-191363

2107

2108

HEMBA1007319	2.84	3.61	6.73	5.21	6.12	3.32		
HEMBA1007320	1.29	1.22	3.12	4.19	3.45	2.42		
HEMBA1007322	19.97	17.81	27.74	45.24	39.42	37.31	**	+
HEMBA1007323	4.54	6.69	11.47	6	6.36	6.21		
HEMBA1007326	4.58	3.85	13.34	8.29	8.07	9		
HEMBA1007327	3.37	3.98	8.91	6.14	9.31	8.98		
HEMBA1007332	3.12	3.47	5.42	5.27	7.56	5.33		
HEMBA1007341	1.4	2.51	3.24	2.93	3.38	3.36		
HEMBA1007342	1.06	2.05	3.02	1.52	3.23	1.98		
HEMBA1007347	3.39	3.24	6.34	4.55	7.34	6.32		
HEMBA1007353	2.43	2.22	4.99	2.68	6.9	3.01		
HEMBA1000005	1.57	2.54	5.35	2.68	4.65	3.64		
HEMBA1000008	2.19	2.53	5.99	3.51	6.31	3.71		
HEMBA1000018	2.21	2.16	7.13	9.9	9.79	7.24		
HEMBA1000024	3.71	2.15	5.4	5.13	5.77	6.39		
HEMBA1000025	2.11	2.09	3.55	1.68	5.13	2.62		
HEMBA1000030	3.12	3.53	6.58	6.62	7.77	6		
HEMBA1000036	5.3	4.76	5.04	6.95	8.19	5.93	*	+
HEMBA1000037	4.43	3.64	4.73	4.63	8.38	5.32		
HEMBA1000039	1.17	0.96	3.98	2.61	4.11	2.95		
HEMBA1000044	1.22	2.35	4.26	5.28	5.58	5.36	*	+
HEMBA1000048	3.2	1.7	3.48	3.99	5.4	3.96		
HEMBA1000050	2.32	1.55	3.33	2.97	3.98	2.85		
HEMBA1000054	2.03	2.08	7.07	4.49	5.09	3.98		
HEMBA1000055	42.59	36.75	92.41	100.33	86.52	89.35		
HEMBA1000059	2.5	2.65	11.34	10.96	11.52	14.73		
HEMBA1000072	6.84	7.77	58.85	73.22	97.61	76.22	*	+
HEMBA1000081	2.85	3.56	10.79	5.69	6.19	7.23		
HEMBA1000083	1.13	1.95	5.38	4.88	5.89	5.33		
HEMBA1000089	1.14	2.53	4.54	6.03	6.73	5.43	*	+
HEMBA1000094	4.12	4.3	8.4	4.24	5.04	6.62		
HEMBA1000097	2.48	1.71	7.91	4.75	4.55	4.3		
HEMBA1000099	2.69	2.07	6.27	5.18	6.75	5.64		
HEMBA1000103	7.19	5.28	18.55	13.99	19.26	16.16		
HEMBA1000106	3.91	3.75	8.15	4.24	6.4	5.9		
HEMBA1000113	1.25	1.54	3.33	1.39	4.14	2.31		
HEMBA1000119	2.19	2.17	5.66	3.34	6.12	4.05		
HEMBA1000133	21.01	22.21	30.57	43.5	66.13	60.69	*	+
HEMBA1000134	4.92	2.95	8.69	13.39	9.79	9.8		
HEMBA1000136	7.14	8.81	29.63	23.11	26.28	28.23		
HEMBA1000141	1.98	2.85	6.18	4.95	5.75	6.16		
HEMBA1000144	2.05	2.59	4.85	3.09	5.36	1.38		
HEMBA1000147	3.77	2.08	4.51	4.55	7.39	3.07		
HEMBA1000152	0.79	1.45	3.42	5.13	5.15	3.45		
HEMBA1000154	0.98	1.11	3.43	2.59	2.89	2.47		
HEMBA1000155	0.88	0.54	3.15	3.33	4.15	1.92		
HEMBA1000173	3.35	3.72	12.14	10.1	10.89	7.51		
HEMBA1000175	1.85	1.32	3.39	6.06	4.09	3.86		
HEMBA1000176	1.48	4.03	6.12	3.43	9.75	5.03		
HEMBA1000198	0.88	1.72	3.64	2.6	3.59	3.22		

(1056)

特開2002-191363

2109

2110

HEM BB1000208	1.12	1.52	3.04	1.74	3.23	2.69		
HEM BB1000209	1.62	1.54	3.76	3.32	3.94	3.28		
HEM BB1000212	1.88	1.03	3.26	4.37	3.93	1.98		
HEM BB1000215	1.61	2.13	3.8	4.67	5.49	5.32	*	+
HEM BB1000217	5.67	4.97	11.23	15.21	18.81	11.78	*	+
HEM BB1000218	2.13	2.28	11.05	7.47	10.07	7.05		
HEM BB1000226	2.63	3.26	6.02	4.02	5.86	3.41		
HEM BB1000230	1.39	1.83	3.73	1.95	4.31	2.58		
HEM BB1000240	6.04	8.26	10.01	3.97	2.7	1.75	*	-
HEM BB1000244	1.39	1.64	3.51	2.51	2.63	1.68		
HEM BB1000250	1.17	0.99	1.12	1.94	1.25	1.12		
HEM BB1000258	1.71	1.94	5.8	4.38	5.63	3.27		
HEM BB1000264	2.49	3.12	11.01	8.64	8.34	8.1		
HEM BB1000266	2.81	2.65	5.52	3.38	5.95	3.71		
HEM BB1000272	4.76	4.16	6.06	8.38	6.88	7.45	*	+
HEM BB1000274	1.51	1.15	3.17	2.54	3.18	1.88		
HEM BB1000276	1.12	1.84	4.72	3.1	4.01	2.43		
HEM BB1000284	0.94	1.81	2.89	2.83	3.11	1.65		
HEM BB1000307	1.52	1.7	4.78	2.8	5.31	3.27		
HEM BB1000309	1.43	2.73	3.07	3.09	3.56	2.19		
HEM BB1000312	1.99	1.38	5.18	7.03	7.2	4.35		
HEM BB1000317	0.17	1.62	3.32	2.6	4.73	2.14		
HEM BB1000318	1.11	2.69	3.85	2.28	4.46	1.68		
HEM BB1000332	3.12	3.84	4.37	3.75	3.95	2.72		
HEM BB1000335	0.77	2.35	4.66	6.16	4.66	3.44		
HEM BB1000336	0.99	1.11	3.59	2.09	3.52	2.29		
HEM BB1000337	4.3	5.06	20.22	22.86	24.84	22.82		
HEM BB1000338	2.11	1.92	5.86	7.13	8.92	4.71		
HEM BB1000339	1.66	1.76	5.84	3.75	4.99	3.84		
HEM BB1000341	1.4	1.91	3.68	3.06	4.77	2.83		
HEM BB1000343	2.51	3.15	6.96	7.24	8.68	7.46		
HEM BB1000354	3.26	3.5	10.36	7	8.93	8.07		
HEM BB1000358	1.09	2.11	3.82	3.43	2.83	1.93		
HEM BB1000369	1.93	2.33	3.87	5.96	6.54	2.89		
HEM BB1000373	1.77	2.73	3.91	2.26	6.2	2.94		
HEM BB1000374	3.27	4.06	9.34	9.58	13.36	6.95		
HEM BB1000376	2.71	3.92	12.28	10.03	6.99	7.71		
HEM BB1000383	60.87	62.14	104.01	69.28	57.52	83.25		
HEM BB1000391	1.8	2.66	4.57	4.89	6.18	4.29		
HEM BB1000399	2.51	3.79	3.69	3.93	5.72	3.71		
HEM BB1000402	1.61	2.06	3.33	2.67	5.3	1.72		
HEM BB1000404	1.34	1.15	5.18	2.56	5.29	1.81		
HEM BB1000407	2.2	3.36	6.76	5.57	5.75	4.66		
HEM BB1000420	1.93	1.46	3.86	4.33	4.76	4.54	*	+
HEM BB1000430	38.77	36.24	61.06	51.76	34.69	50.02		
HEM BB1000434	3.05	4.73	9.02	6.54	6.59	6.63		
HEM BB1000438	1.13	1.83	4.16	2.23	4	1.39		
HEM BB1000441	2.26	3	7.35	5.44	8	4.78		
HEM BB1000447	29.84	32.01	39.91	35.88	44.02	33.55		
HEM BB1000449	1.3	1.31	3.72	1.51	3.04	1.54		

(1057)

特開2002-191363

2111

2112

HEMBB1000453	8.61	8.04	13.39	14.23	18.78	13.74		
HEMBB1000455	1.29	1.97	3.19	3.13	5.46	3.54		
HEMBB1000472	2.3	2.28	4.22	4.07	4.35	3.52		
HEMBB1000480	1.9	3.59	7.03	5.71	6.63	5.87		
HEMBB1000486	2.15	2.98	6.93	4.82	7.86	5.9		
HEMBB1000487	1.21	1.79	4.48	2.66	4.8	2.57		
HEMBB1000490	3.67	4.13	12.61	7.92	8.7	6.55		
HEMBB1000491	1.36	2.91	5	3.83	4.78	4.42		
HEMBB1000492	3.02	4.04	6.84	5.63	6.94	5.34		
HEMBB1000493	1.57	1.71	3.26	2.41	5.38	3.27		
HEMBB1000510	1.32	1.71	4.94	4.4	5.61	4.21		
HEMBB1000516	5.64	7.71	36.22	16.62	18.58	17.09		
HEMBB1000518	0.88	1.22	2.63	2.21	4.27	1.73		
HEMBB1000523	1.32	2.78	7.41	3.33	7.74	4.1		
HEMBB1000530	2.83	2.51	9.72	6.06	7.81	6.64		
HEMBB1000542	3.08	4.55	9.39	8.48	14.07	9.82		
HEMBB1000550	4.84	2.87	4.77	10.48	5.74	5.33		
HEMBB1000554	2.14	2.26	8.65	6.43	11.59	7.19		
HEMBB1000556	2.64	2.68	4.48	3.1	4.67	4.6		
HEMBB1000564	1.81	1.4	5.87	4.26	5.34	6.18		
HEMBB1000567	1.39	1.71	3.7	2.1	3.87	3.3		
HEMBB1000569	3.78	2.72	8.76	4.88	7.13	5.65		
HEMBB1000573	3.48	3.44	11.31	7.33	9.55	7.56		
HEMBB1000575	12.42	2.7	10.57	7.16	8.38	12.35		
HEMBB1000579	2.12	3.75	4.84	3.01	8.61	4.04		
HEMBB1000585	0.83	1.19	3.19	2	4.1	3.35		
HEMBB1000586	2.18	1.41	4.28	4.07	4.23	4.11		
HEMBB1000589	2.98	1.58	3.62	3.38	3.75	3.69		
HEMBB1000591	2.62	2.15	3.96	4.44	5.26	4.64	*	+
HEMBB1000592	2.05	1.2	3.18	3.87	3.39	2.15		
HEMBB1000593	10.25	5.67	51.69	53.87	82.75	77.88	*	+
HEMBB1000595	6.42	5.2	11.24	12.51	17.1	9.55		
HEMBB1000598	1.57	1.69	5.91	2.94	6.76	4.77		
HEMBB1000611	0.94	1.16	2.08	1.15	2.79	1.66		
HEMBB1000617	2.01	3.04	9.31	6.14	8.79	6.97		
HEMBB1000623	2.51	3.08	4.64	5.58	5.83	3.9		
HEMBB1000630	3.23	2.5	3.78	2.42	5.54	2.51		
HEMBB1000631	8.91	10.69	18.75	22.52	23.76	22.55	*	+
HEMBB1000632	6.77	8.77	20.85	27.2	18.4	23.31		
HEMBB1000636	9.52	15.91	22.42	25.26	21.65	19.96		
HEMBB1000637	6.63	9.77	19.44	17.77	24.39	20.28		
HEMBB1000638	1.44	1.41	3.23	3.6	5.29	3.34		
HEMBB1000642	3.47	2.31	7.58	7.65	9.33	9.93		
HEMBB1000643	0.71	1.87	2.71	1.62	3.54	2.12		
HEMBB1000649	2.25	2.22	6.45	4.94	7.61	5.72		
HEMBB1000652	1.8	2.21	5.33	5.13	5.14	4.93		
HEMBB1000655	1.07	1.17	3.24	1.64	4.74	2.01		
HEMBB1000665	0.52	1.08	2.23	1.69	2.92	2.01		
HEMBB1000668	1.85	1.46	2.76	5.07	5.42	4.1	**	+
HEMBB1000671	2.36	2.01	6.77	7.03	7.81	6.94		

(1058)

特開 2002-191363

2113

2114

HEMBB1000673	0.75	1.27	2.92	2.84	4.63	2.43		
HEMBB1000679	3.26	2.84	5.59	4.42	7.19	5.76		
HEMBB1000684	1.83	2.53	6.6	5.01	6.92	5.6		
HEMBB1000692	0.93	2	2.46	1.77	2.5	1.09		
HEMBB1000693	0.96	1.29	2.47	1.6	2.79	1.34		
HEMBB1000705	2.61	2.52	4.85	4.97	8.2	6.53		
HEMBB1000706	0.78	1.07	2.18	2.56	2.93	1.06		
HEMBB1000709	3.53	2.92	8.39	8.16	7.99	10.26		
HEMBB1000714	1.41	2.85	9.32	5.31	10.37	8.79		
HEMBB1000725	1.61	2.22	4.35	3.04	6.22	4.72		
HEMBB1000726	1.88	2.34	8.76	5.63	7.1	4.83		
HEMBB1000729	1.82	3.28	4.3	3.3	5.21	2.79		
HEMBB1000738	1.94	2.6	5.55	3.99	5.53	6.15		
HEMBB1000749	4.06	4.15	7.47	7.48	9.56	8.27		
HEMBB1000763	5.81	5.56	6.21	6.65	9.9	6.61		
HEMBB1000770	2.76	2.06	8.8	7.73	9.62	8.83		
HEMBB1000774	1.62	2.75	3.7	3.07	4.39	2.34		
HEMBB1000777	5.17	5.49	7.42	6.86	4.9	7.37		
HEMBB1000781	3.7	4.19	6.89	7.64	5.28	6.83		
HEMBB1000788	0.87	1.79	2.45	2.65	4.88	1.35		
HEMBB1000789	1.91	2.22	3.01	3.1	6.55	1.86		
HEMBB1000790	1.97	2.15	4.48	4.59	4.21	2.53		
HEMBB1000794	1.46	1.8	2.85	2.97	3.84	2.06		
HEMBB1000807	2.55	2.72	5.18	3.57	4	4.26		
HEMBB1000809	30.31	26.87	132.99	158.22	156.74	195.14	*	+
HEMBB1000810	1.98	2.67	4.51	3.76	6.08	4.12		
HEMBB1000821	1.98	1.93	2.98	2.05	4.45	1.79		
HEMBB1000822	1.08	1.97	2.31	1.65	5.31	1.46		
HEMBB1000826	1.36	1.99	3.57	3.77	6.11	3.6		
HEMBB1000827	2.48	2.89	5.83	2.67	5.05	2.99		
HEMBB1000831	3.4	2.31	5.67	3.84	7.74	2.95		
HEMBB1000835	1.76	1.94	6.2	7.59	7.62	7.47	*	+
HEMBB1000840	1.27	2.95	6.89	4.48	7.19	3.01		
HEMBB1000848	2.08	3.45	5.63	5.39	6.45	5.3		
HEMBB1000852	1.26	2.16	2.8	1.07	4.51	1.55		
HEMBB1000857	7.65	6.49	8.13	7.01	10.69	11.53		
HEMBB1000858	3.7	3.13	7.3	7.07	9.38	7.31		
HEMBB1000867	2.21	1.84	4.9	3.02	5.55	4.04		
HEMBB1000870	1.64	2.37	4.56	2.84	5.31	3.63		
HEMBB1000876	1.48	2.86	3.91	4.54	3.22	3.93		
HEMBB1000881	3.35	5.56	10.5	6.12	5.88	3.85		
HEMBB1000883	1.02	2.68	2.2	3.03	3.32	2.58		
HEMBB1000887	16.9	14.54	43.41	67.39	61.26	59.84	*	+
HEMBB1000888	1.03	1.67	2.39	1.63	3.92	1.86		
HEMBB1000890	2.93	3.36	10.85	6.01	8.62	7.68		
HEMBB1000893	3.28	2.54	5.46	4.5	6.14	5.57		
HEMBB1000900	1.27	1.53	2.98	2.06	2.54	1.58		
HEMBB1000905	5.09	3.75	6.6	10.05	9.45	8.77	**	+
HEMBB1000908	3.34	2.79	3.01	4.48	4.71	5.7	**	+
HEMBB1000910	1.74	2.91	2.55	2.09	3.56	2.24		

(1059)

特開2002-191363

2115

2116

HEMBB1000913	1.41	1.51	2.22	2.8	3.41	1.91		
HEMBB1000915	32.08	25.6	50.05	48	58.92	51.07		
HEMBB1000917	2.1	2.78	5.72	2.99	4.52	3.44		
HEMBB1000927	1.45	1.24	1.82	1.49	3.25	1.88		
HEMBB1000932	0.66	2.06	2.74	1.81	3.41	1.61		
HEMBB1000933	7.47	7.12	10.71	12.88	12.78	19.19		
HEMBB1000936	1.44	1.96	2.87	3.75	6.44	3.55		
HEMBB1000939	7.86	7.14	9.02	15.98	15.3	18.25	**	+
HEMBB1000941	1.53	1.86	3.17	3.99	4.46	3.52	*	+
HEMBB1000947	3.53	3.34	4.61	4.67	6.8	5.72		
HEMBB1000954	1.08	1.82	2.54	1.62	4.01	3.08		
HEMBB1000959	0.77	1.41	3.11	2.23	4.42	1.87		
HEMBB1000973	1.21	1.12	2.6	2.07	4.47	3.2		
HEMBB1000975	0.9	1.28	1.55	2.48	2.86	1.7	*	+
HEMBB1000981	1.54	0.66	1.99	1.84	3.47	2.03		
HEMBB1000985	1.67	1.82	2.71	3.59	3.99	3.24	*	+
HEMBB1000991	0.99	1.35	2.83	1.46	3.75	2.43		
HEMBB1000996	4.89	3.12	6.78	6.75	6.64	11.18		
HEMBB1001000	1.86	1.39	4.06	3.07	5.46	3.88		
HEMBB1001004	1.15	1.32	2.57	2.56	4.55	2.42		
HEMBB1001008	1.48	1.79	2.75	2.44	4.6	2.88		
HEMBB1001011	1.34	1.15	1.53	2.47	2.24	2.9	**	+
HEMBB1001014	1.31	1.43	2.3	2.73	4.84	4.1	*	+
HEMBB1001020	1.17	0.75	2.77	1.77	2.67	2.26		
HEMBB1001024	3.31	1.72	6.27	5.47	7.56	6.82		
HEMBB1001026	5.14	4.03	5.16	5.46	7.67	5.44		
HEMBB1001037	2	1.45	4.73	3.52	5.69	6.67		
HEMBB1001042	0.52	1.15	2.69	1.29	3.61	0.87		
HEMBB1001046	1.18	1.28	2.16	1.67	3.82	0.96		
HEMBB1001047	1.01	1.7	3.79	2.2	3.83	3.83		
HEMBB1001048	2.5	2.34	7.02	4.34	11.02	6.93		
HEMBB1001051	1.44	2.62	3.23	3.95	6.26	3.9		
HEMBB1001056	1.61	2.67	4.89	3.75	5.7	3.78		
HEMBB1001058	1.3	1.92	4.72	2.64	6.92	2.63		
HEMBB1001060	0.69	0.68	1.75	2.05	4.61	1.85		
HEMBB1001063	1.23	1.83	3.52	2.43	4.21	2.9		
HEMBB1001068	1.84	3.62	3.59	3.46	7.14	5.2		
HEMBB1001082	2.24	2.57	5.98	5.38	6.93	6.36		
HEMBB1001095	6.39	7.45	11.76	14.04	14.61	13.16	*	+
HEMBB1001096	1.3	1.91	3.05	3.21	4.12	3.49		
HEMBB1001101	7.41	8.19	9.74	19.33	13.1	16.69	*	+
HEMBB1001102	1.04	1.47	4.57	3.6	6.07	4.46		
HEMBB1001104	1.66	1.89	3.98	3.1	4.87	4.25		
HEMBB1001105	1.57	1.59	2.13	3.11	4.82	2.71		
HEMBB1001112	9.44	8.91	73.3	100.88	136.14	131.28	*	+
HEMBB1001113	2.11	1.94	9.1	5.65	8.02	6.81		
HEMBB1001114	1.88	2.27	5.18	4.16	7.06	4.82		
HEMBB1001115	5.78	7.88	14.52	16.77	9.5	14.78		
HEMBB1001117	1.7	1.52	2.92	1.85	2.79	1.79		
HEMBB1001119	1.69	1.57	4	2.29	3.74	2.28		

(1060)

特開2002-191363

2117

2118

HEMBB1001126	1.85	1.88	3.63	2.69	4.96	2.9		
HEMBB1001133	3.15	2.42	4.24	6.56	6.15	6.44	**	+
HEMBB1001137	1.97	2.2	4.4	3.28	6.56	5.42		
HEMBB1001142	2.96	2.68	10.51	9.2	11.69	10.38		
HEMBB1001145	3.25	3.56	7.39	6.11	7.7	6.59		
HEMBB1001151	5.67	6.58	9.21	12.85	7.67	7.84		
HEMBB1001153	2	1.89	4.57	3.79	5.37	2.9		
HEMBB1001158	6.96	6.74	12.17	12.04	9.95	11.16		
HEMBB1001169	1.71	2.45	4.42	2.89	4.21	2.86		
HEMBB1001170	1	1.85	3.27	1.26	2.5	1.61		
HEMBB1001175	1.43	1.54	5.16	3.39	6.36	4.14		
HEMBB1001177	3.63	2.4	7.54	5.8	6.94	6.69		
HEMBB1001182	2.6	3.69	4.89	3.89	6.54	3.87		
HEMBB1001192	3.3	3.09	16.1	15.27	20.06	15.75		
HEMBB1001199	1.16	2.27	1.91	1.43	3.96	1.15		
HEMBB1001200	1.86	1.66	3.14	2.43	5.38	2.29		
HEMBB1001208	2.02	2.04	4.56	2.96	6.19	2.74		
HEMBB1001209	2.98	2.28	5.75	5.22	6.56	4.92		
HEMBB1001210	5.14	4.28	7.8	11.05	6.08	10.33		
HEMBB1001215	9.57	10.46	17.69	17.91	15.75	16.96		
HEMBB1001217	1.78	2.13	4.39	2.04	3.69	1.89		
HEMBB1001218	4.28	3.37	5.47	4.52	5.98	4.6		
HEMBB1001221	1.72	1.65	2.75	1.54	4.29	1.34		
HEMBB1001224	2.2	2.46	3.81	2.98	6.08	3.7		
HEMBB1001230	1.51	2.09	4.36	2.55	4.95	2.22		
HEMBB1001234	5.24	6.05	29.26	31	44.04	30.75		
HEMBB1001235	12.72	10.54	21.49	13.27	8.47	10.71		
HEMBB1001237	11	10.54	21.03	32.1	26.16	36.86	*	+
HEMBB1001242	4.82	5.68	8.63	6.92	6.97	4.51		
HEMBB1001244	1.08	1.1	3.9	1.47	4.36	1.36		
HEMBB1001249	1.26	1.63	2.99	1.84	5.52	1.98		
HEMBB1001253	1.53	1.92	5.68	1.96	4.89	2.39		
HEMBB1001254	1.27	1.19	3.73	1.22	5.09	2.45		
HEMBB1001266	2	4.32	4.75	4.49	6.09	4.63		
HEMBB1001267	3.51	2.92	9.98	8.43	7.63	8.01		
HEMBB1001271	2.25	2.93	3.89	2.59	5.85	3.91		
HEMBB1001282	2.27	2.68	3.9	2.77	4.86	2.99		
HEMBB1001287	54.06	45.71	83.21	72.79	57.33	77.51		
HEMBB1001288	2.45	2.58	3.64	4.57	6.08	3.31		
HEMBB1001289	4.64	5.82	12.2	6.93	9.11	6.88		
HEMBB1001290	2.82	1.27	4.55	2.89	4.14	1.96		
HEMBB1001294	1.03	1.91	3.2	2.93	3.95	2.51		
HEMBB1001299	7.06	7.64	12.49	16.26	14.41	17.87	*	+
HEMBB1001302	2.16	2.34	2.41	1.75	3.39	1.64		
HEMBB1001304	1.73	1.34	2.6	1.81	5.2	1.67		
HEMBB1001314	1.16	1.07	2.47	1.3	3.63	1.25		
HEMBB1001315	1.25	1.62	1.46	0.87	4	0.9		
HEMBB1001317	2.1	3.38	6.51	4.12	8.01	4.41		
HEMBB1001326	0.88	1.54	2.36	1.51	3	1.69		
HEMBB1001331	2.11	2.79	2.81	3.78	6.14	4.55	*	+

(1061)

特開2002-191363

2119

2120

HEMBB1001335	1.39	0.9	1.44	1.4	2.81	1.92		
HEMBB1001337	1.86	1.7	3.15	3.34	4.72	4.51	*	+
HEMBB1001339	4.17	3.87	5.91	5.83	7.91	5.25		
HEMBB1001344	1.27	1.36	2.25	1.62	3.44	1.29		
HEMBB1001346	2.17	2.32	7.45	5.89	6.23	5.7		
HEMBB1001348	0.68	1.37	4.05	1.38	3.68	2.82		
HEMBB1001350	2.06	2.09	4.17	2.07	7.74	2.57		
HEMBB1001356	1.4	1.9	2.33	1.74	5.54	2.29		
HEMBB1001364	0.53	1.28	1.25	1.49	2.47	1.43		
HEMBB1001366	1.61	1.71	3.46	4.17	4.27	3.82	*	+
HEMBB1001367	1.11	2.19	3.79	5.05	6.25	5.75	*	+
HEMBB1001369	0.56	1.29	2.54	1.95	3.39	2.82		
HEMBB1001380	3.13	3.56	6.65	5.43	7.36	7.75		
HEMBB1001381	8.45	6.07	9.53	10.2	14.39	11.86		
HEMBB1001384	3.48	4.92	5.66	9.52	13.27	11.91	**	+
HEMBB1001387	1.19	1.57	3.1	2.36	4.34	1.33		
HEMBB1001394	1.53	1.3	1.68	2.7	2.82	1.89	*	+
HEMBB1001407	0.68	0.83	0.99	0.62	1.33	0.97		
HEMBB1001410	1.35	1.04	1.78	2.44	2.63	1.74		
HEMBB1001413	1.68	1.84	3.32	3.48	3.27	4.04		
HEMBB1001419	2.56	2.24	4.42	3.61	5.47	4.6		
HEMBB1001421	2.29	1.66	2.18	1.31	3.16	0.95		
HEMBB1001424	0.51	1.2	1.67	-0.1	1.58	0.41		
HEMBB1001426	2.04	1.51	3.7	2.66	5.67	4.21		
HEMBB1001429	7.11	5.76	9.83	22.69	19.97	19.53	**	+
HEMBB1001436	3.13	2.51	6.8	7.5	6.44	7.24		
HEMBB1001443	5.61	6.48	20.67	20.46	27.07	22.15		
HEMBB1001449	2.02	2	4.92	4.26	6.35	4.27		
HEMBB1001454	1.2	1.96	3.77	3.74	5.13	2.99		
HEMBB1001458	4.72	6.48	12.41	7.88	9.04	6.89		
HEMBB1001461	0.55	1.38	2.01	2.11	2.65	1.07		
HEMBB1001463	2.28	2.1	3.7	3.95	5.03	4.66	*	+
HEMBB1001464	1.73	1.29	3.62	2.66	3.92	2.27		
HEMBB1001466	1.15	1.84	2.75	1.88	3.77	2.73		
HEMBB1001482	1.76	2.21	4.36	2.94	5.36	3.97		
HEMBB1001500	1.01	1.08	1.77	1.96	4.39	1.97		
HEMBB1001505	3	3.32	5.87	9.35	14.06	11.17	*	+
HEMBB1001521	2.06	2.43	5.4	5.15	5.78	7.14		
HEMBB1001527	2.63	4.74	11.16	8.69	9.66	9.6		
HEMBB1001530	4.15	3.51	6.57	9.43	12.39	7.05		
HEMBB1001531	1.11	1.34	4.62	2.99	5.16	4.17		
HEMBB1001532	0.63	1.86	2.77	1.86	4.41	2.15		
HEMBB1001535	1.99	2.01	4.22	3.34	4.32	5.75		
HEMBB1001536	2.18	2.65	6.37	4.62	6.87	5.45		
HEMBB1001537	1.31	2.24	3.7	3.21	6.12	2.75		
HEMBB1001542	4.39	4.72	6.28	5.26	7.83	5.7		
HEMBB1001543	7.84	3.58	8.49	8.13	7.08	5.38		
HEMBB1001547	2.02	2.25	2.65	4.2	4.27	2.79	*	+
HEMBB1001548	2.53	2.62	11.82	17.73	29.92	23.34	*	+
HEMBB1001551	0.89	1.7	4.47	2.93	5.96	2.65		

(1062)

特開2002-191363

2121

2122

HEM BB1001555	2.13	2.79	4.78	3.73	5.8	4.64		
HEM BB1001562	1.9	2.64	4.27	2.23	3.46	3		
HEM BB1001564	132.08	140.08	310.28	333.18	233.12	279.03		
HEM BB1001565	1.72	1.97	3.9	3.77	4.68	2.07		
HEM BB1001569	0.79	0.8	3.04	2.49	3.68	1.44		
HEM BB1001573	1.9	1.04	3.58	4.18	5.53	4.12	*	+
HEM BB1001585	1.5	1.96	10.91	3.75	7.14	4.87		
HEM BB1001586	1.53	2.27	3.22	4.27	5.45	1.95		
HEM BB1001588	1.33	2.9	5.74	4.72	6.32	5.06		
HEM BB1001595	2.68	3.33	6.92	3.78	4.84	4.7		
HEM BB1001596	3.4	2.57	3.74	2.67	5.36	2.54		
HEM BB1001599	1.45	1.57	3.21	3.07	3.47	2.06		
HEM BB1001603	1.99	2.45	4.17	5.7	8.16	4.04		
HEM BB1001606	1.35	2.28	2.42	2	2.39	1.53		
HEM BB1001612	4.31	3.07	9.25	8.81	8.09	8.9		
HEM BB1001618	1.53	1.62	3.86	2.84	4.48	2.31		
HEM BB1001619	2.11	3.03	3.92	5.71	5.1	4.37	*	+
HEM BB1001623	2.21	2.38	3.16	2.16	5.5	3.37		
HEM BB1001625	3.73	3.04	4.33	2.93	4.81	3.79		
HEM BB1001630	1.31	2.36	3.54	1.84	4.3	1.23		
HEM BB1001635	1.78	1.64	3.76	2.08	3.32	1.34		
HEM BB1001637	1.76	1.14	3.98	2.09	4.4	3.58		
HEM BB1001641	1.43	1.68	2.78	3.17	3.73	2		
HEM BB1001653	2.18	3.17	5.61	3.96	6.63	3.5		
HEM BB1001665	1.08	2.17	2.04	2.5	4.43	0.88		
HEM BB1001666	2.14	1.95	3.52	2.45	4.88	1.79		
HEM BB1001667	2.37	2.25	3.26	2.94	5.13	3.17		
HEM BB1001668	3.19	2.11	5.15	2.45	6.42	2.69		
HEM BB1001669	0.98	2.02	3.19	1.04	4.53	1.38		
HEM BB1001670	4.02	4.82	6.88	10.7	9.71	8.65	*	+
HEM BB1001673	1.48	2.97	3.61	3.51	4.52	4.43		
HEM BB1001675	1.83	3.27	4.65	4.68	5.78	4.88		
HEM BB1001679	2.52	2.34	5.06	2.19	3.87	1.88		
HEM BB1001684	2.13	1.55	3.89	5.17	6.77	5.05	*	+
HEM BB1001685	3.41	1.61	4.43	2.91	6.24	2.49		
HEM BB1001695	1.9	2.22	4.43	1.38	3.88	2.12		
HEM BB1001703	1.25	2.3	5.74	3.58	3.79	4.1		
HEM BB1001704	1.39	2.16	4.58	4.23	5.02	3.94		
HEM BB1001706	2.76	2.6	3.58	5.6	6.26	4.87	**	+
HEM BB1001707	1.35	2.01	2.87	2.25	3.67	2.8		
HEM BB1001717	1.68	2.21	3.23	2.61	3.34	2.83		
HEM BB1001731	13.81	13.48	24.03	11.02	23.09	25		
HEM BB1001734	3.47	3.35	7.62	6.88	9.22	4.18		
HEM BB1001735	1.35	1.4	3.4	1.58	3.52	2.03		
HEM BB1001736	5.01	6.14	7.87	7.15	10.91	8.11		
HEM BB1001747	0.92	1	3.23	1.87	3.67	2.82		
HEM BB1001749	4.71	2.99	9.39	7.29	5.99	8.16		
HEM BB1001753	3.79	3.3	5.5	7.4	8.97	9.3	**	+
HEM BB1001756	0.53	2.05	1.89	2.31	3.91	2.73		
HEM BB1001757	1.08	1.8	2.64	3.04	4.86	4.54	*	+

(1063)

特開2002-191363

2123

2124

HEMBB1001760	1.32	0.98	3.74	1.49	3.56	2.13		
HEMBB1001762	0.9	0.61	2.62	1.57	2.95	2.07		
HEMBB1001780	9.82	12.28	11.34	16.64	26.06	22.06	*	+
HEMBB1001785	0.89	1.24	1.02	0.62	2.88	1.64		
HEMBB1001788	3.22	1.26	5.17	5.76	6.13	5.3		
HEMBB1001793	5.6	4.73	18.12	22.08	20.38	22.86	*	+
HEMBB1001797	1.61	1.82	2.28	2.94	4.97	5.54	*	+
HEMBB1001802	13.28	9.91	67.77	85.35	82.8	81.27	*	+
HEMBB1001812	2.32	2.58	6.49	7.68	8.28	9.74	*	+
HEMBB1001815	128.22	114.78	102.97	87.37	89.21	62.14	*	-
HEMBB1001816	1.95	2.19	3.97	3.73	6.31	5.73		
HEMBB1001831	0.69	0.98	1.54	0.72	3.28	1.16		
HEMBB1001834	16.15	9.68	133.91	102.49	173.48	141.04		
HEMBB1001836	4.07	1.99	11.4	5.89	6.97	6.7		
HEMBB1001839	0.89	0.86	1.43	1.08	1.95	1.21		
HEMBB1001841	80.32	59.68	120.73	35.74	39.04	23.13	*	-
HEMBB1001844	5.26	4.72	9.73	10.15	10.68	8.11		
HEMBB1001847	6.93	4.24	8.6	9.06	14.77	10.3		
HEMBB1001848	25.33	21.68	40.92	49.73	73.94	62.17	*	+
HEMBB1001850	3.07	2.93	5.25	3.75	7.19	3.95		
HEMBB1001859	13.4	8.82	20.85	39.61	27.01	40.75	*	+
HEMBB1001863	1.7	3.65	7.66	6.89	8.88	7.5		
HEMBB1001867	1.69	1.93	3.16	3.16	4.14	3		
HEMBB1001868	2.15	1.53	2.56	1.31	3.57	1.28		
HEMBB1001869	1.5	2.3	4.23	4.62	8.15	3.58		
HEMBB1001872	1.21	0.79	2.23	1.77	4.17	1.75		
HEMBB1001874	1.92	1.2	2.07	2.58	2.47	3.03	*	+
HEMBB1001875	0.83	1.96	1.7	2.05	3.53	1.37		
HEMBB1001880	2.68	2.17	6.79	4.41	8.91	8.35		
HEMBB1001899	0.6	1.62	2.15	2.26	4.68	1.91		
HEMBB1001903	5.56	5.68	9.61	6.06	6.65	6.7		
HEMBB1001905	2.04	2.82	5.49	4.98	6.07	7.14		
HEMBB1001906	0.67	1.65	2.64	2.22	5.14	2.62		
HEMBB1001908	1.82	1.63	5.26	3.31	6.56	3.59		
HEMBB1001910	1.92	1.16	2.64	3.2	4.41	4.61	*	+
HEMBB1001911	2.06	1.11	4.2	5.73	4.81	4.33		
HEMBB1001915	2.53	3.01	5.61	7.13	7.84	9.4	*	+
HEMBB1001921	2.19	2.21	7.12	6.56	8.47	8.25		
HEMBB1001922	1.74	1.77	3.66	4.91	5.85	3.7		
HEMBB1001925	1.48	2.24	4.57	3.07	4.73	3.39		
HEMBB1001930	0.46	0.94	1.67	0.91	3.18	1.01		
HEMBB1001944	1.72	1.88	5.45	4.83	4.6	4.49		
HEMBB1001945	2.15	1.05	3.58	2.25	3.97	2.12		
HEMBB1001947	2.28	1.13	3.23	5.35	5.98	3.15		
HEMBB1001950	3.49	1.95	5.38	5.15	5.46	3.3		
HEMBB1001952	1.41	2.05	4.72	2.31	4.73	3.1		
HEMBB1001953	1.62	2.09	3.45	3.63	5.78	3.75		
HEMBB1001957	1.16	2.02	4.28	2.68	4.19	2.19		
HEMBB1001959	2.05	3.31	4.07	4.22	5.75	3.68		
HEMBB1001962	5.54	2.61	5.11	3.34	6.28	2.54		

(1064)

特開2002-191363

2125

2126

HEMBB1001967	2.59	2.46	5.62	6.11	8.49	6.15		
HEMBB1001973	2.25	2.4	6.14	7.62	10	7.28	*	+
HEMBB1001978	2.08	1.71	6.29	7.2	7.57	5.83		
HEMBB1001983	9.23	8.69	24.64	38.93	34.91	36.79	*	+
HEMBB1001987	1.78	2.34	3.64	1.66	4.75	2.38		
HEMBB1001988	2.02	1.92	3.42	2.92	5.17	1.85		
HEMBB1001990	7.65	7.72	9.18	12.44	11.53	15.42	*	+
HEMBB1001996	1.54	1.47	3.89	1.61	3.51	1.22		
HEMBB1001997	1.46	2.25	6.1	4.2	5.98	4.23		
HEMBB1001999	10.91	11.08	16.84	24.47	26.58	22.28	**	+
HEMBB1002002	1.08	1.58	3.52	1.91	2.76	2.39		
HEMBB1002005	1.88	2.91	4.8	4.82	7.6	4.22		
HEMBB1002009	2.32	2.48	3.03	2.24	6.23	2.7		
HEMBB1002013	0.96	2.07	3.78	1.95	4.26	1.41		
HEMBB1002015	3.95	4.25	9.47	5.82	8.92	6.73		
HEMBB1002024	45.16	34.47	111.32	113.31	106.76	120.55		
HEMBB1002035	2.15	1.91	2.87	2.11	4.5	2.68		
HEMBB1002039	1.18	2.29	5.1	3.28	5.9	2.98		
HEMBB1002041	3.31	4.13	8.49	15.49	14.42	13.38	**	+
HEMBB1002042	3.97	4.66	9.49	8.09	10.63	9.94		
HEMBB1002043	1.34	2.21	4.61	5.97	5.24	3.36		
HEMBB1002044	0.4	1.19	2.68	1.25	4.19	1.92		
HEMBB1002045	2.83	2.5	10.03	6.34	7.63	4.56		
HEMBB1002049	1.31	1.4	3.77	1.71	4.36	1.73		
HEMBB1002050	1.62	1.61	4.5	3.31	4.53	2.94		
HEMBB1002051	1.17	1.13	2.9	2.59	5.05	4.37		
HEMBB1002068	1.69	2.44	2.43	2.3	4.42	2.07		
HEMBB1002069	3.39	3.94	7.83	6.86	7.55	5		
HEMBB1002075	0.72	1.94	3.33	2.99	3.52	2.37		
HEMBB1002079	1.2	1.8	1.89	1.22	2.84	1.3		
HEMBB1002080	1.74	1.85	4.78	1.55	6.02	2.41		
HEMBB1002082	1.03	1.85	4.59	2.38	4.38	1.96		
HEMBB1002084	25.86	22.68	51.44	33.52	35.54	38.37		
HEMBB1002088	13.92	15.78	22.14	29.46	37.25	35.66	**	+
HEMBB1002092	2.51	2.24	4.48	5.34	3.65	6.27		
HEMBB1002094	3.21	2.62	8.2	5.72	7.27	6.04		
HEMBB1002103	2.42	2.97	3.51	3.74	5.58	4.47		
HEMBB1002109	4.27	3.47	4.84	5.72	7.83	6.36	*	+
HEMBB1002115	42.37	37.4	91.88	95.86	101.94	101.65		
HEMBB1002120	0.89	1.22	2.91	0.86	2.94	2.41		
HEMBB1002121	0.75	1.56	1.63	1.5	4.66	2.53		
HEMBB1002134	11.99	11.22	112.59	98.93	166.1	133.77		
HEMBB1002136	1.29	1.65	2.9	2.59	3.26	2.58		
HEMBB1002138	10.48	9.64	20.72	18.78	23.06	19.4		
HEMBB1002139	1.84	1.6	5.46	4.69	6	5.84		
HEMBB1002141	1.53	0.83	3.44	1.48	4.64	2.54		
HEMBB1002142	1.85	2	4.95	3.22	5.98	6.03		
HEMBB1002145	1.62	0.83	2.96	1.49	3.07	2.42		
HEMBB1002152	1.27	1.19	3.15	2.32	6.36	3.41		
HEMBB1002162	1.25	1.55	3.92	3.42	5.14	3.61		

(1065)

特開 2002-191363

2127

2128

HEM BB1002173	4.18	1.09	5.58	2.77	4.48	3.84		
HEM BB1002189	2.78	1.95	6.14	7.01	8.25	5.93		
HEM BB1002190	1.81	2.2	6.36	8.01	6.93	8.36	*	+
HEM BB1002193	1.84	1.06	2.06	4.53	4.37	5.48	**	+
HEM BB1002217	3.82	2.26	6.02	3.61	6.06	3.23		
HEM BB1002218	3.91	3.3	7.58	4.94	6.15	5.68		
HEM BB1002228	2.28	2.9	6.17	6.68	7.94	6.97		
HEM BB1002232	1.15	1.4	2.24	2.14	5.79	3.03		
HEM BB1002245	0.86	0.84	2.34	1.53	2.47	1.05		
HEM BB1002247	1.72	0.59	2.44	1.38	2.24	1.03		
HEM BB1002249	2.65	1.64	3.1	3.39	4.08	4.01	*	+
HEM BB1002254	1.35	1.35	3.83	2.64	3.27	2.86		
HEM BB1002255	0.99	1.37	2.6	1.19	2.93	1.4		
HEM BB1002266	1.33	0.83	2.07	0.73	1.99	0.54		
HEM BB1002271	14.89	9.5	29.42	32.9	41.53	37.28	*	+
HEM BB1002280	1.62	0.78	1.6	1.55	2.93	1.12		
HEM BB1002296	11.83	12.31	18.29	31.87	21.06	23.09	*	+
HEM BB1002300	0.78	2.31	3.48	1.49	4.63	3.12		
HEM BB1002302	1.17	2.17	3.26	2.49	4.76	3.04		
HEM BB1002306	1.83	1.96	4.28	2.91	4.34	2.6		
HEM BB1002316	0.66	1.38	2.36	1.19	2.9	0.97		
HEM BB1002326	0.93	1.68	4.52	4.35	4.41	3.9		
HEM BB1002327	0.99	0.99	2.66	1.46	2.95	2.06		
HEM BB1002329	2.89	3	3.81	6.39	5.88	5.53	**	+
HEM BB1002340	0.6	1.8	2.05	2.29	3.38	2.22		
HEM BB1002342	8.12	9.23	14.09	21.68	18.15	18.03	*	+
HEM BB1002358	1.09	3.22	6.37	6.52	7.91	9.2		
HEM BB1002359	1.09	2.55	4.29	4.66	5.4	3.54		
HEM BB1002364	1.28	1.82	2.33	3.17	5.15	3.11		
HEM BB1002366	13.63	21.17	32.35	56.28	57.48	53.09	**	+
HEM BB1002371	0.83	0.63	1.72	2.32	2.82	2.1	*	+
HEM BB1002381	0.97	1.16	1.74	2.83	3.16	6.26		
HEM BB1002383	1.07	3.17	4.18	3.1	4.5	4.37		
HEM BB1002387	0.98	2.36	2.68	2.09	3.68	2.96		
HEM BB1002409	6.85	7.27	46.98	69.94	70.04	64.2	*	+
HEM BB1002413	3.92	2.99	8.34	9.46	8.16	10.16		
HEM BB1002415	0.84	1.28	2.79	2.36	3.49	2.11		
HEM BB1002424	1.04	1.17	1.63	2.89	3.04	2.66	**	+
HEM BB1002425	1.12	1.69	5.86	6.46	10.1	5.26		
HEM BB1002427	1.5	1.59	2.32	3.72	7.82	3.65		
HEM BB1002442	2.29	1.57	4.33	4.99	8.58	8.89	*	+
HEM BB1002447	2.61	2.7	5.56	6.1	6.6	5.63		
HEM BB1002453	2.5	2.48	6.56	6.31	7.55	5.25		
HEM BB1002457	1.54	2.08	4.77	3.69	4.61	4.8		
HEM BB1002458	0.48	1.53	2.5	2.2	2.35	1.66		
HEM BB1002463	1.36	1.84	6.55	6.24	6.11	8.87		
HEM BB1002465	1.12	1.18	2.4	2.86	2.25	1.59		
HEM BB1002477	0.71	0.66	4.43	3	4.39	4.86		
HEM BB1002479	22.08	21.58	27.54	16.12	19.41	17.27	*	-
HEM BB1002489	0.86	3.02	3.9	5.73	5.51	7.68	*	+

(1066)

特開2002-191363

2129

2130

HEM BB1002492	1.27	1.23	3.07	3.53	4.08	3.39	*	+
HEM BB1002495	1.85	1.85	3.2	2.61	5.02	3.98		
HEM BB1002502	0.94	2.52	2.81	1.77	4.83	3.27		
HEM BB1002509	0.73	1.8	2.65	2.03	2.43	1.27		
HEM BB1002510	0.49	1.68	3.06	1.78	2.5	0.81		
HEM BB1002520	1.46	2.47	5.44	6.62	7.57	8.61	*	+
HEM BB1002522	0.82	1.88	4.42	2.31	6.8	2.07		
HEM BB1002527	11.47	13.79	12.46	24.19	10.37	17.52		
HEM BB1002530	1.43	2.15	3.44	2.93	4.92	2.26		
HEM BB1002531	0.46	1.32	2.04	1.23	2.99	0.35		
HEM BB1002534	1.35	2.27	2.73	4.54	4.08	3.92	**	+
HEM BB1002536	6.58	5.93	46.38	45.93	63.71	42.88		
HEM BB1002544	3.91	3.45	6.89	6.79	7.87	7.99		
HEM BB1002545	0.92	2.76	2.83	3.21	4.15	4.29		
HEM BB1002550	1.32	1.69	1.86	2.99	4.68	2.42		
HEM BB1002556	2.9	3.54	9.69	8.73	8.12	10.62		
HEM BB1002571	17.25	14.03	19.8	21.91	16.59	24.61		
HEM BB1002579	3.32	2.05	4.87	4.38	6.6	6.39		
HEM BB1002582	1.79	2.11	5.59	5.77	6.47	5.63		
HEM BB1002584	2.82	1.94	6.09	3.94	4.67	3.62		
HEM BB1002587	6.39	5.82	10.63	11.3	9.04	9.94		
HEM BB1002590	1.6	3.07	7.46	5.86	7.3	5.84		
HEM BB1002596	1.5	2.01	3.17	5.59	4.94	4.21	*	+
HEM BB1002600	1.55	2.72	3.81	5.02	7.18	3.93		
HEM BB1002601	1.28	2.23	3.9	2.51	5.59	2.99		
HEM BB1002603	2.37	1.64	5.48	3.53	6.59	5.6		
HEM BB1002607	1.48	1.15	4.34	2.59	4.26	2.99		
HEM BB1002610	1.2	0.96	3.48	1.95	3.79	3.45		
HEM BB1002613	0.96	2.41	4.31	3.98	5.39	3.72		
HEM BB1002614	3.18	3.34	5.35	3.87	6.08	9.76		
HEM BB1002615	1.47	3.29	4.63	2.45	3.83	2.67		
HEM BB1002617	0.67	3.09	2.88	2.1	3.34	3.4		
HEM BB1002623	2.31	3.63	4.36	3.96	6.28	5.91		
HEM BB1002624	2.7	1.56	7.52	7.96	8.72	8.3		
HEM BB1002631	1.65	2	4.28	2.14	3.72	1.74		
HEM BB1002635	1.84	1.74	3.55	3.31	3.64	3.56		
HEM BB1002644	7.22	9.04	15.98	18.52	23.94	22.55	*	+
HEM BB1002654	5.22	4.21	7.77	7.92	7.33	11.74		
HEM BB1002661	1.93	2.16	3.96	1.99	4.13	5.33		
HEM BB1002663	1.59	1.8	3.85	5.45	5.58	5.17	*	+
HEM BB1002664	1.28	2.4	4.43	5.05	9.22	6.76	*	+
HEM BB1002677	1.88	1.83	1.86	1.81	4.79	2.34		
HEM BB1002683	2.68	2.21	9.21	5.67	5.9	8.45		
HEM BB1002684	1.71	0.81	2.53	1.92	2.74	2.63		
HEM BB1002686	1.23	1.39	2.88	1.45	3.37	1.64		
HEM BB1002692	0.99	1.4	1.87	2.5	2.53	2.98	*	+
HEM BB1002693	1.75	1.75	4.12	5.03	5.74	3.46		
HEM BB1002697	1.09	2.8	2.73	4.17	5.58	5.34	*	+
HEM BB1002699	1.59	2.27	4.93	4.72	6.74	6.97		
HEM BB1002702	1.63	1.5	2.54	1.76	3.25	3.33		

(1067)

特開2002-191363

2131

2132

HEMBB1002705	4.2	2.84	6.79	8.83	8.26	7.92	*	+
HEMBB1002712	8.55	1.32	2.38	2.92	4.06	1.4		
IMR321000028	1.03	1.71	2.88	1.63	2.76	1.63		
IMR321000031	1.71	2.59	3.51	5.86	4.35	5.31	*	+
IMR321000034	21.95	15.41	30.37	33.73	19.59	34.65		
IMR321000039	5.81	7.11	14.41	14.72	15.71	13.99		
IMR321000044	0.81	2.37	1.44	1.01	3.26	2.06		
IMR321000063	79.52	80.12	127.61	224.23	199.69	128.8		
IMR321000085	21.02	18.07	26.38	30.28	48.13	47.89	*	+
IMR321000089	1.51	1.42	3.86	3	6.7	5.84		
IMR321000091	4.79	2.91	6.5	8.35	11.38	8.55	*	+
LIVER1000004	8.04	9.67	34.15	55.9	56.53	48.86	*	+
LIVER1000008	1.13	1.36	3.06	1.68	4	2.17		
LIVER1000011	3.03	5.9	26.65	37.8	54.37	45.77	*	+
LIVER1000022	2.75	3.66	7.75	9.39	9.82	9.17	*	+
LIVER1000025	1.78	2.77	5.47	9.83	10.83	7.7	*	+
LIVER1000030	1.05	0.96	2.12	2.04	2.56	1.23		
LIVER1000045	1.33	1.37	3.11	5.11	5.12	5.89	**	+
LIVER1000046	1.01	1.53	3.86	4.14	7.82	5.34		
LIVER1000072	1.61	1.26	5.23	12.42	9.54	12.21	**	+
LIVER1000077	0.33	1.79	1.97	1.87	2.84	3.14		
LIVER1000080	1.53	3	5.81	5.96	4.24	5.41		
LIVER1000086	6.38	7.69	47.4	69.84	79.87	70.57	*	+
LIVER1000092	1.6	1.46	3.09	3.85	3.83	2.41		
LIVER1000095	0.91	2.31	2.56	2.16	2.46	1.55		
LIVER1000097	1.26	0.74	2.49	2.18	2.84	2.25		
LIVER1000098	0.43	1.37	2.57	2.76	3.95	2.29		
LIVER1000100	3.3	2.82	5.82	4.99	7.44	3.74		
LIVER1000101	0.36	1.81	2.4	1.69	4.25	2.74		
LIVER1000106	0.83	1.95	1.79	0.97	2.8	1.29		
LIVER1000108	1.36	2.93	4.31	4.35	4.12	3.43		
LIVER1000115	1.12	1.57	4.32	6.38	6.13	6.9	*	+
LIVER1000120	1.45	0.95	2.23	1.39	1.46	0.73		
LIVER1000138	0.6	1.27	1.86	2.11	2.36	1.6		
LIVER1000146	1.38	2.69	6.24	7.17	7.01	7.02		
LIVER1000148	0.88	1.24	2.65	1.68	5.68	3.51		
LIVER1000157	30.11	26.71	67.66	123.41	85.61	124.96	*	+
LIVER1000161	1.3	1.59	2.3	1.73	3.37	1.84		
LIVER1000167	3.07	3.63	14.08	20.36	22.31	21.82	*	+
LIVER1000174	1.53	1.68	1.84	2.1	3.43	1.29		
LIVER1000185	2.42	2.55	5.16	4.37	4.72	4.23		
LIVER1000187	0.96	1.55	4.84	6.64	4.17	3.5		
LIVER1000190	3.77	3.48	5.95	4.71	6.24	4.47		
LIVER1000192	2.37	2.92	3.93	4.1	5.23	4.04		
MAMMA1000009	1.39	2.55	5.12	3.62	4.72	3.46		
MAMMA1000015	1.72	1.59	4.78	5.23	4.42	6.42		
MAMMA1000019	0.69	2.48	3.4	4.27	4.81	3.1		
MAMMA1000020	2.79	2.35	5.63	6	7.86	5.75		
MAMMA1000024	0.65	1.76	3.79	2.42	2.91	1.61		
MAMMA1000025	1.92	2.56	6.92	4.96	6.72	5.6		

(1068)

特開2002-191363

2133

2134

MAMMA1000043	1.06	2.36	6.43	6.93	8.22	6.6		
MAMMA1000045	1.38	2.01	4.84	2.68	3.96	2.89		
MAMMA1000046	1.74	2.44	3.18	2.88	4.5	2.37		
MAMMA1000055	8.51	8.71	9.57	9.38	10.74	9.36		
MAMMA1000057	4.4	3.29	7.56	8.38	9.78	8.16		
MAMMA1000060	26.78	24.33	45.25	48.69	33.84	48.6		
MAMMA1000069	2.13	1.65	4.1	3.43	3.14	2.41		
MAMMA1000084	2.88	3	5.81	5.64	8.15	7.51		
MAMMA1000085	2.75	3.74	7.02	6.45	5.82	6.97		
MAMMA1000092	1.45	2.97	3.8	4.64	5.15	4.55	*	+
MAMMA1000096	4.45	4.96	9.29	8.15	9.11	6.09		
MAMMA1000097	2.4	2.96	3.86	5.93	6.01	6.97	**	+
MAMMA1000102	1.94	1.59	4.27	4.25	6.16	4.44		
MAMMA1000103	1.52	1.65	5	2.39	4.56	2.83		
MAMMA1000106	1.25	2.15	5.1	2.3	4.48	3.27		
MAMMA1000117	1.19	2.12	3.72	1.84	3.32	2.77		
MAMMA1000118	1.03	2.06	3.08	3.38	3.21	4.56		
MAMMA1000129	1.06	2.1	2.97	1.73	2.7	1.31		
MAMMA1000133	1.09	1.96	3.67	2.8	3.87	2.02		
MAMMA1000134	1.23	2.08	4.28	2.27	4.61	1.93		
MAMMA1000139	1.45	1.91	2.69	2.13	3.98	1.85		
MAMMA1000141	1.97	2.27	5.47	3.67	3.61	3.42		
MAMMA1000143	1.66	1.1	2.55	3.83	3.52	1.74		
MAMMA1000150	4.11	4.95	8.99	6.49	6.4	8.66		
MAMMA1000155	1.87	2.71	4.35	5.46	5.69	6.41	*	+
MAMMA1000163	1.65	2.82	2.62	3.45	4.54	4.97	*	+
MAMMA1000171	1.96	2.43	5.53	4.12	6.52	4.97		
MAMMA1000173	3.5	5.27	10.33	17.47	17.44	18.26	**	+
MAMMA1000175	1.58	1.8	2.89	4.06	5.14	3.7	*	+
MAMMA1000183	1.14	2.12	5.29	3.92	4.7	4.91		
MAMMA1000191	3.25	3.34	14.68	12.56	14.7	18.23		
MAMMA1000192	5.76	8.53	9.67	16.66	19.91	17.65	**	+
MAMMA1000193	1.68	1.54	0.86	1.33	2.25	1.94		
MAMMA1000198	1.88	1.99	5.53	4.44	6.49	5.57		
MAMMA1000204	1.75	2.25	3.39	3.56	4.13	2.85		
MAMMA1000207	1	3.2	3.41	2.86	4.96	3.47		
MAMMA1000214	1.76	2.08	3.68	2.84	4.42	4.74		
MAMMA1000220	6.19	6.12	11.61	12.49	18.06	16.72	*	+
MAMMA1000221	0.57	1.04	1.68	1.14	4.51	0.87		
MAMMA1000226	0.48	1.06	2.07	1.49	3.19	1.88		
MAMMA1000227	0.93	1.23	1.6	2.73	3.67	3.46	**	+
MAMMA1000230	1	1.23	1.77	2.38	3.04	2.94	**	+
MAMMA1000241	2.9	2.2	4.19	7.24	5.8	7.61	**	+
MAMMA1000245	76.63	70.15	118.95	141.45	166.09	104.88		
MAMMA1000248	6.79	4.17	13.48	13.18	13.44	18.8		
MAMMA1000251	1.68	1.72	4.7	5.55	5.39	5.29		
MAMMA1000254	1.24	1.22	3.59	2.14	5.61	5.02		
MAMMA1000257	5.39	2.62	25.06	32.2	43.78	35.79	*	+
MAMMA1000262	15.48	9.75	18.2	40.81	33.23	34.89	**	+
MAMMA1000264	0.99	1.2	2.3	4.43	2.57	3.4	*	+

(1069)

特開2002-191363

2135

2136

MAMMA1000266	1.25	0.79	2.73	4.21	5.33	4.03	*	+
MAMMA1000270	2.43	1.94	4.57	6.16	7.16	7.58	*	+
MAMMA1000271	6.01	3.26	8.54	8.94	6.17	8.1		
MAMMA1000277	0.89	0.93	2.56	2.46	2.75	2.09		
MAMMA1000278	1.84	2.01	4.29	2.18	5.06	3.51		
MAMMA1000279	1.82	1.74	4.33	3.51	5.72	4.35		
MAMMA1000283	0.99	1.51	2.36	1.37	2.66	2.71		
MAMMA1000284	2.65	2.51	8.31	6.28	8.49	8.01		
MAMMA1000287	1.58	2.13	6.27	5.55	6.94	7.1		
MAMMA1000294	4.72	5.45	9.44	3.84	8.21	4.74		
MAMMA1000298	0.87	1.36	2.51	1.55	3.1	0.95		
MAMMA1000302	0.9	1.18	4.73	2.22	4.9	2.56		
MAMMA1000303	0.92	1.62	2.63	4.16	4.06	3.22	*	+
MAMMA1000305	1.07	1.28	2.73	2.47	2.74	2.1		
MAMMA1000307	2.29	3.03	9.61	15.85	14.04	14.38	*	+
MAMMA1000309	0.57	1.61	3.69	4.6	3.65	4.61		
MAMMA1000312	3.55	4.99	8.08	6.19	5.18	6.55		
MAMMA1000313	1.06	2.31	2.34	1.79	3.98	3.43		
MAMMA1000331	1.08	1.65	3	3.33	6.01	3.21		
MAMMA1000335	7.38	9.1	14.27	19.49	16.28	16.92	*	+
MAMMA1000339	0.33	0.39	2.17	1.46	2.09	0.66		
MAMMA1000340	1.43	1.33	4.12	4.37	2.72	2.15		
MAMMA1000348	1.2	1.27	4.6	3.14	4.82	6.11		
MAMMA1000356	1.93	2.21	4.93	3.08	5.42	6.29		
MAMMA1000358	2.93	3.97	5.02	7.42	7.47	4.9		
MAMMA1000360	1.41	1.92	4.6	3.76	5.2	4.63		
MAMMA1000361	2.2	3.45	8.9	8.94	10.67	8.93		
MAMMA1000363	1.09	1.69	3.86	1.87	4.37	4.1		
MAMMA1000370	0.92	0.71	1.76	2.02	2.62	2.57	*	+
MAMMA1000371	2.09	1.73	6.35	10.02	12.1	10.1	*	+
MAMMA1000372	4.45	4.1	12.88	12.01	12.92	11.97		
MAMMA1000385	1.79	2.36	6.41	6.41	7.66	8.72		
MAMMA1000388	1.93	3.02	6.03	4.7	4.53	5.06		
MAMMA1000395	1.3	2.46	3.12	1.69	3.49	0.8		
MAMMA1000402	1.69	1.68	5.62	3.33	4.35	4.63		
MAMMA1000403	1.7	2.36	5.05	5.45	5.81	3.96		
MAMMA1000410	0.87	1.25	2.71	3.23	3.35	3.25	*	+
MAMMA1000413	1.52	0.47	2.48	3.51	3.76	3.61	*	+
MAMMA1000414	1.08	1.53	3.03	2.94	4.91	1.81		
MAMMA1000416	3.3	4.01	10.2	15.8	23.14	20.47	*	+
MAMMA1000421	2.61	2.83	6.11	7.7	7.42	7.09	*	+
MAMMA1000422	2.83	2.53	7.46	9.18	6.64	12.05		
MAMMA1000423	1.7	1.26	6	5.9	6.62	5.89		
MAMMA1000424	0.88	1.7	3.17	1.91	2.38	1.07		
MAMMA1000429	8.73	10.07	13.78	14.98	16.3	11.17		
MAMMA1000431	1.6	1.27	4.27	5.22	6.32	4.26		
MAMMA1000432	1.05	2.33	2.85	2.63	2.82	1.41		
MAMMA1000437	4.61	4.75	8.44	10.54	11.52	8.12		
MAMMA1000444	2.53	4.15	8.55	7.55	10.17	10.13		
MAMMA1000446	1.19	2.07	3.87	2.03	3.63	2.49		

(1070)

特開2002-191363

2137

2138

MAMMA1000449	1.77	1.59	3.54	3.37	4.31	3.22		
MAMMA1000457	4.44	4.82	7.12	7.2	6.88	6.22		
MAMMA1000458	1.27	2.22	4.83	2.52	4.03	1.94		
MAMMA1000468	0.55	1.12	2.2	0.51	2.25	1.16		
MAMMA1000472	1.15	2.3	4.42	4.77	6.36	5.79	*	+
MAMMA1000473	1.95	1.72	3.59	3.45	5.46	3.17		
MAMMA1000477	3.86	3.29	5.67	8.71	9.92	7.97	**	+
MAMMA1000478	2.85	3.26	7.41	5.76	9.1	7.57		
MAMMA1000483	4.16	3.16	8.3	8.09	6.5	8.63		
MAMMA1000490	1.65	2.61	3.68	2.66	4.96	2.14		
MAMMA1000496	1.18	1.7	3.44	1.3	3.79	2.01		
MAMMA1000500	0.68	1.79	3.22	1.41	3.2	2.86		
MAMMA1000501	3.04	3.89	7.86	13.71	15.02	12.51	**	+
MAMMA1000503	0.84	2.08	2.21	3.52	3.52	2.27		
MAMMA1000506	10.14	8.79	32.66	34.77	26.7	18.31		
MAMMA1000510	3.24	3.5	4.59	10.97	10.76	13.61	**	+
MAMMA1000515	2.12	1.54	4.56	5.97	7.55	6.2	*	+
MAMMA1000516	2.18	2.4	6.29	3.89	3.85	5.07		
MAMMA1000522	1.04	1.47	4.39	2.4	3.68	1.85		
MAMMA1000524	2.04	2.09	3.53	3.82	6.18	3.96		
MAMMA1000528	3.74	2.72	2.05	2.7	4.09	2.88		
MAMMA1000534	0.91	2.35	2.11	1.86	2.91	2.04		
MAMMA1000541	2.85	3.16	11.29	11.22	8.81	12.04		
MAMMA1000550	1.21	2.73	1.86	2.46	6.65	4.64		
MAMMA1000556	1.78	1.32	4.25	2.66	3.37	1.4		
MAMMA1000559	1.32	1.49	5.56	2.92	4.2	3.46		
MAMMA1000565	1.82	2.74	3.93	2.13	4.18	4.22		
MAMMA1000567	0.99	2.16	3.77	2.3	4.07	3.64		
MAMMA1000576	3.72	3.12	13.12	10.45	9.76	10.02		
MAMMA1000582	2.07	2.7	5.64	4.13	4.81	6.31		
MAMMA1000583	1.16	2.33	2.45	2.19	4.47	2.93		
MAMMA1000585	1.66	2.04	4.19	3.82	5.38	3.23		
MAMMA1000587	1.64	1.51	3.73	3.12	5.07	3.49		
MAMMA1000591	0.96	1.34	3.11	1.45	3.74	1.72		
MAMMA1000594	2.3	1.76	4.92	3.55	7.68	5.16		
MAMMA1000597	4.42	3.09	9.64	9.46	9.63	10.35		
MAMMA1000605	2.84	3.94	11.44	18.34	15.85	17.89	*	+
MAMMA1000612	1.91	2.15	5.22	3.85	4.33	4.95		
MAMMA1000614	3.11	2.71	9.4	7.48	6.07	6.34		
MAMMA1000616	1.66	1.79	2.44	2.1	5.09	3.45		
MAMMA1000621	1.39	1.67	3.36	3.15	6.31	7.02		
MAMMA1000623	1.08	1.04	3.83	0.92	2.66	2.3		
MAMMA1000625	7.39	6.32	23.76	19.68	25.39	29.8		
MAMMA1000635	0.89	0.68	1.61	0.76	1.75	0.64		
MAMMA1000643	1.47	1.11	1.94	4.21	3.77	6.82	*	+
MAMMA1000646	4.68	3.61	9.55	17.22	16.4	16.44	**	+
MAMMA1000652	1.98	1.61	3.56	4.21	5.34	5.24	*	+
MAMMA1000657	2.28	1.78	4.51	2.18	4.48	3.26		
MAMMA1000664	1.78	1.49	5.35	2.9	6.43	6.85		
MAMMA1000667	1.24	1.68	2.17	1.96	5.64	2.41		

(1071)

特開 2002-191363

2139

2140

MAMMA1000668	0.71	1.11	3.23	2.44	3.76	1.74		
MAMMA1000669	0.76	0.97	2.01	0.9	3.94	2.06		
MAMMA1000670	3.27	2.78	4.47	10.4	6.73	9.28	*	+
MAMMA1000672	1.71	3.23	6.88	5.43	5.63	6.03		
MAMMA1000681	0.98	1.19	2.53	1.98	3.73	2.45		
MAMMA1000684	6.87	11.61	18.54	30.76	32.53	30.62	**	+
MAMMA1000696	1.64	3.39	4.99	7.89	14.39	8.69	*	+
MAMMA1000702	3.12	3.07	5.9	7.47	10.05	7.55	*	+
MAMMA1000706	0.63	1.07	1.79	1.08	1.52	0.66		
MAMMA1000707	0.74	1.26	1.76	0.83	1.87	0.63		
MAMMA1000713	1.53	2.14	5.33	5.43	5.8	6.96		
MAMMA1000714	1.19	1.84	4.31	2.64	4.96	4.94		
MAMMA1000718	1.32	2.79	4.84	5.53	7.12	4.37		
MAMMA1000720	1.33	2.19	5.14	4.95	8.51	5.44		
MAMMA1000723	1.22	1.65	4.17	3.26	4.81	3.68		
MAMMA1000731	1.24	1.17	3.11	3.04	4.99	3.26		
MAMMA1000732	1.37	1.59	3.02	4.86	6.5	6.05	**	+
MAMMA1000733	0.58	0.82	1.54	2.31	2.41	1.22		
MAMMA1000734	12.22	11.56	22.62	21.95	22.19	13.18		
MAMMA1000736	4.26	4.34	11.96	4.92	5.77	6.14		
MAMMA1000738	0.8	2.06	3.82	2.52	4.15	1.95		
MAMMA1000744	1.12	2	5.52	3.27	3.97	4.67		
MAMMA1000746	1	2.03	2.24	2.38	4.27	2.48		
MAMMA1000748	8.23	8.93	13.13	15.53	16.06	15.05	*	+
MAMMA1000751	10.46	7.63	32.43	45.16	40.03	54.65	*	+
MAMMA1000752	1.5	2.37	8.68	11.52	14.2	12.68	*	+
MAMMA1000757	1.89	2.48	5.54	7.9	8.53	10.86	*	+
MAMMA1000760	3	2.99	6.77	4.65	8.01	7.12		
MAMMA1000761	1.86	2.58	5.73	4.87	6.29	4.99		
MAMMA1000775	1.37	1.83	4.43	3.23	4.29	2.64		
MAMMA1000776	2.37	2.36	6.3	6.57	7	6.12		
MAMMA1000778	2.14	2.28	5.19	4.95	3.99	3.84		
MAMMA1000781	1.33	1.33	3.06	2.82	2.86	1.23		
MAMMA1000782	1.94	2.36	3.88	2.93	3.11	3.01		
MAMMA1000784	1.28	1.58	3.94	2.22	7.2	3.64		
MAMMA1000788	3.05	4.31	4.38	4.16	5.12	2.21		
MAMMA1000798	1.01	2.86	2.77	2.03	3.56	2.1		
MAMMA1000802	4.36	3.71	8.23	17.49	15.39	19.2	**	+
MAMMA1000810	3.91	4.98	14.15	17.58	19.04	16.15	*	+
MAMMA1000813	1.63	2.11	3.73	2.82	3.15	2.06		
MAMMA1000814	2.56	2.97	7.82	7.28	7.26	6.61		
MAMMA1000824	12.58	11.27	34.16	62.44	72.28	50.62	*	+
MAMMA1000827	1.83	2.04	5.05	3.3	4.77	4.31		
MAMMA1000831	1.45	2.81	2.84	2.55	5.43	1.85		
MAMMA1000838	6.85	9.27	7.86	13.01	8.43	14.52		
MAMMA1000839	4.89	4.41	9	9.02	8.66	14.04		
MAMMA1000841	1.33	2.02	3.11	2.23	4.44	2.29		
MAMMA1000842	2.48	1.88	3.59	3.03	3.74	1.82		
MAMMA1000843	1.26	2.19	3.88	2.26	4.43	2.35		
MAMMA1000845	0.83	1.01	2.4	2.35	3.88	1.24		

(1072)

特開 2002-191363

2141

2142

MAMMA1000851	1.3	3.42	5.35	7.83	5.9	6.65		
MAMMA1000854	2.77	3.7	6.33	5.08	5.68	5.13		
MAMMA1000855	0.37	2.97	2.62	2.51	3.74	1.78		
MAMMA1000856	0.87	1.39	3.11	2.05	6.37	3.19		
MAMMA1000859	9.88	8.56	20.5	19.52	18.47	24.31		
MAMMA1000862	1.13	1.55	3.53	1.17	3.18	0.79		
MAMMA1000863	2.62	2.08	4.72	2.59	5.24	4.91		
MAMMA1000865	0.35	0.82	2.48	0.4	1.84	0.35		
MAMMA1000867	1.08	2.83	2.87	1.95	3.68	4.76		
MAMMA1000875	0.89	2.72	2.34	3.31	3.57	2.59		
MAMMA1000876	1.23	1.64	4.59	2.4	3.37	3.22		
MAMMA1000877	3.15	2.89	9.22	8.39	10.32	10.07		
MAMMA1000878	3.05	3.61	8.94	6.33	8.52	9.62		
MAMMA1000880	1.46	1.15	4.77	2.45	3.96	3.01		
MAMMA1000881	1.81	2.09	4.52	3.77	6.07	4.73		
MAMMA1000883	0.57	0.79	2.1	1.37	2.14	2.11		
MAMMA1000897	0.76	2.39	0.36	1.19	2.99	6.19		
MAMMA1000898	1.06	1.99	1.75	1.41	2.49	3.7		
MAMMA1000905	1.8	2.75	4.68	8.32	6.86	10.64	*	+
MAMMA1000906	1.17	2.49	2.63	2.45	4.27	4.08		
MAMMA1000908	1.59	1.63	4.3	1.77	3.05	1.43		
MAMMA1000911	4.97	6.25	8.37	21.77	20.01	20.98	**	+
MAMMA1000914	1.14	0.85	2.41	1.1	2.23	1.61		
MAMMA1000920	1.99	2.17	4.41	10.82	10.67	9.11	**	+
MAMMA1000921	1.03	1.02	2.41	3.47	3.84	3.17	*	+
MAMMA1000931	2.68	3.44	3.95	6.78	7.66	7.32	**	+
MAMMA1000940	1.67	1.84	6.05	5.78	6.84	6.3		
MAMMA1000941	3.74	2.55	8.61	9.01	10.11	9.46		
MAMMA1000942	2.75	1.85	7.46	8.27	8.47	7.2		
MAMMA1000943	2.16	2.84	10.49	9.04	11.08	8.74		
MAMMA1000952	2.6	1.93	8.65	9.03	7.47	6.75		
MAMMA1000956	0.93	1.24	3.11	3.64	3.1	3.47		
MAMMA1000957	2.5	1.41	2.62	4.81	5.73	6.85	**	+
MAMMA1000962	3.25	3.57	10.48	13.62	11.18	16.85	*	+
MAMMA1000966	1.85	2.19	6.04	6.34	6.23	7.04		
MAMMA1000968	1.6	1.46	5.49	4.79	5.62	4.97		
MAMMA1000972	2.4	1.41	3.83	3.34	3.91	4.32		
MAMMA1000973	5.14	3.37	12.58	7.02	8.56	10.31		
MAMMA1000975	1.44	1.99	3.34	2.37	4.9	5.05		
MAMMA1000976	2.46	2.71	8.57	9.22	11.17	8.92		
MAMMA1000979	1.46	2.62	3.06	4.34	4.41	7.71		
MAMMA1000986	5.75	5.32	10.24	8.83	11.32	13.95		
MAMMA1000987	1.44	1.36	3.99	2.43	3.74	5.66		
MAMMA1000988	3.76	4.86	8.88	10.18	11.34	10.5	*	+
MAMMA1000994	9.82	7.58	15.88	12.02	11.56	9.25		
MAMMA1000998	1.51	1.07	3.13	4.21	5.42	4.04	*	+
MAMMA1001003	1.98	1.83	5.97	4.2	6.86	4.39		
MAMMA1001007	0.38	1.03	1.77	0.14	1.38	0.32		
MAMMA1001008	11.76	11.09	40.52	56.73	50.93	45.37	*	+
MAMMA1001013	3.62	4.16	12.14	8.42	11.54	10.08		

(1073)

特開2002-191363

2143

2144

MAMMA1001014	1.4	1.79	5.49	4.36	5.16	4.26		
MAMMA1001021	0.49	2.08	7.85	6.46	5.47	4.26		
MAMMA1001024	0.85	1.59	3.14	2.55	3.92	1.96		
MAMMA1001025	1.03	1.47	2.94	1.27	2.95	1.41		
MAMMA1001028	1.3	1.09	2.23	3.07	4.73	3.28	*	+
MAMMA1001030	1.63	0.48	2.22	3.06	3.24	2.08		
MAMMA1001035	2.48	2.92	10.31	9.29	11.92	11.54		
MAMMA1001036	4.69	4.01	10.7	11.23	8.22	11.33		
MAMMA1001037	1.91	2.88	6.49	5.28	7.53	4.69		
MAMMA1001038	1.18	1.59	4.28	4.12	3.85	4.45		
MAMMA1001041	1.64	1.87	3.18	3.75	4.04	2.64		
MAMMA1001043	1.09	1.24	3.67	3.44	3.35	3.18		
MAMMA1001050	1.52	1.55	5.94	8.4	8.28	6.51	*	+
MAMMA1001054	2.04	2.58	6.99	9.29	11.07	8.36	*	+
MAMMA1001059	2.66	4.71	9.73	8.85	9.72	8.46		
MAMMA1001066	3.64	2.97	12.26	14.08	12.09	8.96		
MAMMA1001067	1.26	2	4.77	3.53	5.33	3.29		
MAMMA1001072	1.44	2.06	7.76	6.38	7.13	7.2		
MAMMA1001073	1.17	0.79	1.47	1.49	2.74	1.34		
MAMMA1001074	0.78	1.47	3.97	6.24	5.4	4.9	*	+
MAMMA1001075	4.87	4.41	10.48	11.24	9.82	8.21		
MAMMA1001078	1.7	1.83	7.96	9.69	9.99	9.37	*	+
MAMMA1001080	3.77	3.93	6.97	8.71	10.33	5.57		
MAMMA1001082	1.51	2.03	4.03	1.73	5.23	1.6		
MAMMA1001091	1.17	1.36	2.02	1.81	3.03	1.47		
MAMMA1001092	1.93	2.09	4.81	3.17	3.57	1.89		
MAMMA1001094	1.73	4.28	4.65	4	5.62	4.7		
MAMMA1001105	2.45	2.62	7.7	6.99	7.57	6.06		
MAMMA1001110	0.4	1.01	2.74	1.42	2.53	0.47		
MAMMA1001126	1.96	3.09	10.92	8.39	9.27	6.08		
MAMMA1001133	2.5	3.44	10.94	9.48	10.83	10.68		
MAMMA1001139	87.88	86.18	214.31	193.19	47.2	160.92		
MAMMA1001141	1.33	2.89	3.65	3.69	5.25	5.46		
MAMMA1001143	2.02	1.79	4.23	3.95	6.69	4.34		
MAMMA1001145	3.1	2.22	3.39	6.37	7.13	2.84		
MAMMA1001150	1.34	2.48	3.95	3.06	3.31	2.11		
MAMMA1001154	2.16	2.8	5.57	5.44	7.13	5.22		
MAMMA1001159	4.19	4.01	11.06	11.31	5.89	9.45		
MAMMA1001161	4.3	5.27	19.53	18.34	10.8	14.8		
MAMMA1001162	1.98	1.77	3.16	5.25	5.13	2.25		
MAMMA1001181	2.44	2.28	4.87	5.06	4.74	3.62		
MAMMA1001186	2	2.66	4.66	5.38	5.48	3.9		
MAMMA1001189	2.23	3.68	7.17	11	11.17	9.9	*	+
MAMMA1001191	2.54	2.07	5.49	4.37	3.89	2.97		
MAMMA1001198	368.47	416.05	784.82	647.17	738.61	605.52		
MAMMA1001202	11.78	11.85	30.06	34.39	28.74	25.16		
MAMMA1001203	2.57	3.01	7.15	8.72	6.26	5.56		
MAMMA1001206	1.91	3.28	4.5	3.69	6.66	2.65		
MAMMA1001208	2.66	2.93	3.31	3.82	4.95	3.19		
MAMMA1001215	2.9	3.08	6.55	3.49	8.09	4.74		

(1074)

特開2002-191363

2145

2146

MAMMA1001220	2.63	3.03	7.25	7.16	7.17	6.03		
MAMMA1001222	1.25	1.18	4.18	2.18	5.85	0.53		
MAMMA1001223	2.48	3.32	6.53	4.95	6.51	4.1		
MAMMA1001232	2.82	4.27	8.08	12.22	8.82	9.57		
MAMMA1001234	1.25	3.27	3.17	5.05	3.91	3.26		
MAMMA1001237	1.22	1.56	4.21	1.94	3.66	2.09		
MAMMA1001243	2.18	2.28	4.06	4.05	4.89	1.99		
MAMMA1001244	1.22	1.16	2.86	2.96	4.79	2.22		
MAMMA1001249	2.3	1.89	5.93	5.19	5.8	3.75		
MAMMA1001256	3	3.09	8.29	5.89	7.83	8.01		
MAMMA1001259	4.38	3.25	7.15	7.94	9.24	6.63		
MAMMA1001260	1.76	2.71	5.42	6.51	5.33	7.33		
MAMMA1001262	2.1	4.11	5.28	7.86	8.04	6.25	*	+
MAMMA1001268	2	2.16	4.59	2.56	4.23	2.48		
MAMMA1001271	4.84	5.78	17.37	18.29	14.24	15.67		
MAMMA1001274	2.88	3.06	6.17	6.22	8.55	7.93		
MAMMA1001280	2.09	1.48	4.36	1.84	3.78	1.73		
MAMMA1001283	1.63	1.71	6.34	6.88	5.63	4.83		
MAMMA1001284	2.27	2	8.67	5.08	9.09	9.51		
MAMMA1001286	13.83	9.72	17.39	12.15	11.83	14.63		
MAMMA1001289	17.63	13.49	23.32	21.02	26.39	36.8		
MAMMA1001292	3	3.01	5.94	7.26	6.31	6.85	*	+
MAMMA1001296	3.55	3.76	12.61	14.11	12.37	12.8		
MAMMA1001298	1.26	1.7	6.26	4.25	6.78	4.07		
MAMMA1001305	0.86	1.59	4.43	2.49	4.07	2.63		
MAMMA1001309	0.61	0.9	2.7	1.84	3	1.49		
MAMMA1001310	1.72	2.17	3.64	4.81	7.38	4.42		
MAMMA1001322	0.99	1.54	1.83	2.83	1.77	2.13		
MAMMA1001324	1.3	1.12	3.16	2.03	2.83	1.94		
MAMMA1001330	3.35	2.65	9.53	7.93	9.75	5.36		
MAMMA1001333	3.1	3.74	10.23	9.88	11.4	9.07		
MAMMA1001334	5.53	4.17	4.83	10.97	8.23	10.16	**	+
MAMMA1001337	2.49	3.54	6.6	6.99	9.16	8.05	*	+
MAMMA1001341	1.21	1.14	3.48	1.54	5.66	1.41		
MAMMA1001343	2.37	1.89	8.07	8.17	9.75	10.95		
MAMMA1001344	9.59	9.07	11.75	13.63	11.67	15.98		
MAMMA1001346	1.34	1.25	3.9	2.05	3.9	2.94		
MAMMA1001383	3.07	3.61	8.52	8.3	9.02	9.38		
MAMMA1001388	1.62	1.93	5.34	3.38	6.11	4.58		
MAMMA1001396	4.2	2.12	8.12	11.39	10.42	8.68		
MAMMA1001397	2.59	2.27	5.79	8.33	8.96	7.78	*	+
MAMMA1001401	26.87	16.48	32.72	43.47	57.55	45.66	*	+
MAMMA1001408	1.06	1.06	2.57	0.65	4.22	1.19		
MAMMA1001411	1.65	1.26	3.84	4.38	3.33	3.51		
MAMMA1001414	3.12	3.85	5.74	12.58	10.67	15.28	**	+
MAMMA1001415	2.45	3.16	11.93	14.57	20.15	13.69	*	+
MAMMA1001418	0.66	2.2	5.36	3.57	6.04	4.46		
MAMMA1001419	0.8	2.43	4.93	6.03	7.01	3.92		
MAMMA1001420	0.96	3.09	4.5	3.23	4.11	3.41		
MAMMA1001426	20.24	32.21	42.42	44.31	39.63	38.75		

(1075)

特開2002-191363

2147

2148

MAMMA1001428	1.94	2.83	6.35	3.8	6.93	4.33	
MAMMA1001432	1.19	2.33	8.19	5.62	6.19	6.68	
MAMMA1001435	1.43	0.78	3.32	3.48	3.67	2.64	
MAMMA1001442	1.96	3.94	7.41	8.18	8.6	6.63	
MAMMA1001446	2.17	2.57	6.71	6.9	7.34	7.97	
MAMMA1001450	1.22	2.05	3.58	2.81	4.18	2.39	
MAMMA1001452	1.99	1.78	5.92	8.38	6.19	4.83	
MAMMA1001465	3.93	3.25	13.61	16.65	14.6	13.82	
MAMMA1001476	1.63	1.09	4.25	5.87	5.95	4.64	*
MAMMA1001478	2.28	2.12	5.98	3.55	6.27	4.19	+
MAMMA1001479	3.11	4.71	8.32	5.58	6.74	6.21	
MAMMA1001487	1.1	1.14	3.84	4.73	3.26	2.08	
MAMMA1001498	1.93	3.41	7.78	6.17	7.45	5.64	
MAMMA1001501	0.88	1.97	4.49	2.8	4.77	2.36	
MAMMA1001502	1.82	1.91	6.48	3.29	6.29	6.26	
MAMMA1001510	0.48	0.78	2.92	0.54	3.04	1.19	
MAMMA1001522	1.03	1.29	3.94	5.05	4.9	3.39	
MAMMA1001529	0.72	2.06	3.22	3.74	4.07	2.57	
MAMMA1001532	1.74	1.86	4.27	3.79	5.71	3.12	
MAMMA1001533	0.61	1.31	2.9	1.52	3.06	1.64	
MAMMA1001534	0.44	2.59	2.4	1.48	3.64	1.14	
MAMMA1001535	1.38	1.91	3.99	2.12	3.98	2.38	
MAMMA1001547	2.8	2.89	7.77	9.23	8.22	6.22	
MAMMA1001551	1.1	1.48	4.46	2.23	2.88	2.99	
MAMMA1001569	1.27	1.68	3.41	2.03	3.41	1.94	
MAMMA1001575	1.48	2.41	3.42	4.01	4.43	2.81	
MAMMA1001576	4.79	8.23	9.65	14.75	9.39	17.03	
MAMMA1001584	0.89	2.48	3.33	3.11	4	3.09	
MAMMA1001586	1.43	2.41	3.34	3.78	3.31	1.84	
MAMMA1001590	2.96	2.53	5.55	5.44	6.47	6.04	
MAMMA1001599	4.64	7.15	16.79	15.8	15.18	15.06	
MAMMA1001600	1.45	2.22	4.73	2.98	4.68	2.11	
MAMMA1001604	1.03	1.76	3.62	2.35	4.01	1.64	
MAMMA1001606	1.64	2.04	5.15	3.58	5.45	4.27	
MAMMA1001609	1.31	2.37	4.36	3.05	5.43	1.59	
MAMMA1001614	2.91	3.57	6.15	5.94	6.11	4.14	
MAMMA1001615	3.98	2.61	10.12	8.87	8.29	8.41	
MAMMA1001619	7.73	7.8	14.29	16.33	12.93	14.61	
MAMMA1001620	2.53	2.41	7.98	5.77	7.13	4.54	
MAMMA1001623	4.11	4.58	9.3	7.34	9.28	6.75	
MAMMA1001626	0.83	1.98	2.52	3.24	3.93	1.93	
MAMMA1001627	1.11	1.98	3.57	2.63	3.68	1.63	
MAMMA1001630	2.02	3.08	7.83	7.49	7.53	4.29	
MAMMA1001633	2.64	3.12	8.8	12.09	9.59	6.16	
MAMMA1001634	2.83	2.7	6.11	8.69	8.27	6.9	*
MAMMA1001635	5.65	2.39	9.52	7.92	8.3	8.37	+
MAMMA1001649	1.61	1.63	4.71	2.95	4.62	2.53	
MAMMA1001654	8.14	9.45	39	43.4	55	46.79	
MAMMA1001660	19.61	17.92	37.43	40.94	27.03	34	
MAMMA1001663	1.9	4.73	9.42	9.59	9.56	6.58	

(1076)

特開2002-191363

2149

2150

MAMMA1001670	1.12	2.66	3.97	3.65	4.09	2.62		
MAMMA1001671	1.08	1.42	3.56	1.37	4.64	1.77		
MAMMA1001679	6.85	6.37	13.89	11.48	17.04	13.91		
MAMMA1001683	2.15	3.29	9.6	6.58	6.53	6.96		
MAMMA1001686	1.25	1.34	3.77	1.39	2.97	3.06		
MAMMA1001688	113.39	113.61	245.56	392.2	458.41	413	**	+
MAMMA1001689	1.01	3.76	4.1	5.04	3.79	4.44		
MAMMA1001692	1.97	2.59	5.37	3.66	5.3	3.88		
MAMMA1001711	1.99	3.64	8.65	4.35	5.51	6.1		
MAMMA1001715	1.31	1.64	3.95	4.64	4.87	4.13		
MAMMA1001730	2.01	2.15	2.5	2.8	4.42	2.83		
MAMMA1001735	44.73	48.32	102.35	94.99	156.23	119.88		
MAMMA1001740	0.64	1.6	4.59	2.06	3.91	1.95		
MAMMA1001743	9.84	11.15	33.16	41.97	51.62	49.6	*	+
MAMMA1001744	0.63	0.72	0.86	1.1	1.72	1.71	*	+
MAMMA1001745	1.41	2.15	6.15	3.27	4.46	3.93		
MAMMA1001751	1.38	2.41	3.24	2.85	4.51	4.32		
MAMMA1001752	4.7	4.78	9.75	6.12	9.61	8.4		
MAMMA1001754	7.25	7.89	7.34	11.04	9.63	9.39	*	+
MAMMA1001757	1.21	1.1	2.32	2.21	3.25	2.43		
MAMMA1001760	3.87	4.52	20.01	22.91	24.2	27.59	*	+
MAMMA1001764	2.62	2.36	5.97	7.13	10.17	6.51		
MAMMA1001767	1.22	1.55	2.13	1.61	2.96	1.55		
MAMMA1001768	0.57	1.18	4.25	4.74	4.72	4.37		
MAMMA1001769	2.48	2.83	9.22	9.3	9.81	8.94		
MAMMA1001771	2.66	1.58	3.74	2.86	5.85	6.77		
MAMMA1001773	2.7	3.53	3.87	4	6.29	7.61		
MAMMA1001778	0.88	1.92	3.14	3.13	4.21	3.61		
MAMMA1001783	2.01	2.1	11.25	11.63	18.46	13.04		
MAMMA1001785	3	3.52	8.85	10.56	13.38	11	*	+
MAMMA1001788	0.49	0.86	1.21	0.72	1.72	1.11		
MAMMA1001790	1.68	1.67	5.1	2.37	3.73	3.93		
MAMMA1001800	0.83	0.99	1.47	1.5	2.24	3.25		
MAMMA1001804	1.02	1.41	3.18	2.37	4.16	2.4		
MAMMA1001806	2.13	2.78	6.4	3.15	5.5	4.72		
MAMMA1001812	1.46	1.33	5.52	4.21	5.86	5.05		
MAMMA1001815	0.33	1.76	3.07	1.22	3.67	1.24		
MAMMA1001817	3.19	3.38	9.5	6.78	10.89	13.3		
MAMMA1001818	1.68	2.08	3.41	3.94	8.52	3.41		
MAMMA1001819	2.57	4.12	5.82	8.7	10.29	7.87	*	+
MAMMA1001820	2.68	4.51	8.27	7.51	10.98	6.07		
MAMMA1001824	1.66	2.83	8.36	7.55	9.8	7.11		
MAMMA1001832	6.72	7.99	11.85	20.17	21.28	17.21	**	+
MAMMA1001836	1.74	1.66	5.08	4.79	8.19	4.88		
MAMMA1001837	2.61	2	5.84	7.1	9.19	5.37		
MAMMA1001848	1.02	1.61	3.3	2.81	5.33	3.18		
MAMMA1001850	3.79	4.51	9.31	9.98	9.93	14.19		
MAMMA1001851	1.49	2.33	4.98	4.97	4.12	4.02		
MAMMA1001852	2.98	4	9.68	6.4	7.56	6.8		
MAMMA1001854	2.56	3.11	9.16	10.59	10.64	9.98		

(1077)

特開2002-191363

2151

2152

MAMMA1001858	3.11	2.22	5.28	9.93	7.91	8.87	**	+
MAMMA1001864	1.69	1.91	4.09	8.91	6.18	4.37		
MAMMA1001868	0.71	0.92	2.64	1.68	2.58	0.91		
MAMMA1001874	1.2	0.87	2.52	1.06	3.48	1.17		
MAMMA1001878	3.1	3.46	10.86	7.7	13.37	6.77		
MAMMA1001880	2.67	2.99	7.24	5.58	7.17	8.12		
MAMMA1001885	1.14	1.93	6.19	4.7	5.54	4.58		
MAMMA1001890	3.54	3.95	12.93	13.59	13.29	12.2		
MAMMA1001893	3.74	3.42	6.25	6.59	5.49	5.58		
MAMMA1001901	1.13	1.5	5.4	4.53	5.72	2.67		
MAMMA1001907	2.57	1.62	6.43	4.15	7.36	6.34		
MAMMA1001908	3.2	3.36	8.35	11.83	12.96	12.46	*	+
MAMMA1001919	0.23	0.97	3.3	2.24	3.9	2.07		
MAMMA1001931	0.76	1.65	4.04	3.36	5.89	3.25		
MAMMA1001937	2.27	3.15	5.5	6.44	5.06	3.78		
MAMMA1001951	1.74	2.57	6.47	6.48	6.15	4.83		
MAMMA1001956	3.02	3.48	9.72	8.52	7.66	6.76		
MAMMA1001957	3.39	3.51	9.15	7.88	9.47	7.66		
MAMMA1001960	3.1	3.34	7.24	12.06	9.14	6.1		
MAMMA1001963	0.57	0.78	2.14	1.3	2.36	1.06		
MAMMA1001969	1.7	3.43	10.86	8.54	11.14	8.74		
MAMMA1001970	2.86	3.04	8.48	13.11	6.59	6.64		
MAMMA1001978	0.57	1.85	1.76	2.42	3.87	1.53		
MAMMA1001992	2.07	2.04	5.65	6.79	6.75	5.09		
MAMMA1001994	7.97	3.65	11	18.83	13.23	17.17	*	+
MAMMA1002008	3.28	3.77	6.42	3.43	4.06	1.24		
MAMMA1002009	1.46	2.94	5.17	5.73	7.57	4.06		
MAMMA1002011	1.77	1.71	4.26	6.5	6.45	3.37		
MAMMA1002022	1.51	2.1	5.92	6.64	7.42	5.2		
MAMMA1002024	9.79	9.67	19.03	17.61	16.96	22.43		
MAMMA1002032	2.78	2.41	7.25	5.29	6.16	8.07		
MAMMA1002033	3.23	3.95	7.73	11.24	7.23	6.62		
MAMMA1002041	2.87	2.25	3.18	4.74	5.39	1.71		
MAMMA1002042	2.54	2.34	5.66	5.65	5.78	3.76		
MAMMA1002045	2.33	3.51	7.28	8.39	5.05	4.44		
MAMMA1002047	2.58	2.98	8.83	8.7	8.9	6.89		
MAMMA1002056	2.01	5.78	11.14	11.35	10.64	9.14		
MAMMA1002058	1.67	2.61	8.19	4.84	4.66	4.27		
MAMMA1002060	1.08	2.08	1.41	2.5	4.09	1.2		
MAMMA1002065	1.81	2.75	6.04	7.19	5.19	3.26		
MAMMA1002068	2.43	1.84	5.29	4.98	5.6	4.47		
MAMMA1002070	4.5	2.92	4.15	2.58	5.23	2.81		
MAMMA1002078	1.32	1.43	2.94	1.12	4.4	1.07		
MAMMA1002080	7.98	9.71	13.38	14.92	20.84	14.26		
MAMMA1002082	2.54	4.96	13.04	9.67	8.15	7.78		
MAMMA1002084	1.78	3.47	3.38	4.68	4.48	3.6		
MAMMA1002087	1.12	2.15	5.37	3.6	4.67	2.36		
MAMMA1002091	3.79	3.22	4.32	7.18	6.76	6.41	**	+
MAMMA1002093	0.72	1.4	4.31	2.74	4.39	2.33		
MAMMA1002095	2.4	3.22	7.5	4.73	7.52	4.54		

(1078)

特開2002-191363

2153

2154

MAMMA1002108	1.84	1.02	2.63	1.87	3.24	1.31		
MAMMA1002112	2.94	3.4	7.03	12.79	16.02	11.28	**	+
MAMMA1002118	1.02	1.61	2.24	1.41	3.18	2.01		
MAMMA1002119	0.76	2.15	3.61	1.51	3.12	2.54		
MAMMA1002125	1.79	2.61	6.95	4.52	4.19	4.11		
MAMMA1002126	3.72	4.25	9.79	10.08	9.02	11.03		
MAMMA1002128	0.9	2.36	3.07	2.7	3.49	2.88		
MAMMA1002132	3.78	3.24	11.42	6.18	9.05	6.81		
MAMMA1002140	1.46	1.87	3.68	2.18	3.24	2.33		
MAMMA1002142	3.13	3.43	7.06	5.18	7.62	5.46		
MAMMA1002143	5.42	2.27	7.96	7.98	9.87	13.23		
MAMMA1002145	1.47	1.34	3.3	2.9	4.02	2.64		
MAMMA1002147	0.81	1.59	2.9	2.71	4.4	3		
MAMMA1002153	0.99	1.92	5.55	3.52	6.41	4.75		
MAMMA1002155	2.11	1.93	6.76	4.4	6.46	4.46		
MAMMA1002156	0.81	0.8	1.94	0.67	2.63	0.78		
MAMMA1002158	1.38	1.83	5.12	4.09	7.73	5.2		
MAMMA1002164	2.01	2.09	5.86	3.17	3.18	4.04		
MAMMA1002165	4.04	4.29	7.25	8.65	8.1	6.81		
MAMMA1002170	1.01	1.48	154.53	2.65	3.24	4.11		
MAMMA1002174	1.66	2.9	5.88	4.55	7.78	8.58		
MAMMA1002175	3.27	3.3	7.02	6.95	6.64	7.22		
MAMMA1002180	8.59	6.53	35.97	55.49	48.49	51.08	*	+
MAMMA1002198	3.11	2.3	9.33	7.6	11.22	7.13		
MAMMA1002205	2.93	1.66	6.15	6.3	8.04	7.54		
MAMMA1002206	4.6	3.59	8.14	12.4	13.97	11.74	**	+
MAMMA1002209	1.7	1.93	4.03	4.43	4.23	4.57		
MAMMA1002215	4.17	2.72	15.2	11.05	12.43	17.14		
MAMMA1002219	1.57	1.96	4.99	4.84	6.34	5.96		
MAMMA1002224	3.18	2.9	8.18	5.49	7.25	5.86		
MAMMA1002229	3.74	2.21	8.83	8.48	9.26	6.82		
MAMMA1002230	2.02	2.21	6.63	5.31	8.91	6.58		
MAMMA1002233	3.01	1.6	6.08	4.21	7.91	6.14		
MAMMA1002234	3.05	3.06	6.7	8.6	10.45	10.76	*	+
MAMMA1002236	4.13	3.68	14.08	26.56	20.38	24.71	*	+
MAMMA1002243	0.97	2.48	3.48	3.28	3.43	2.96		
MAMMA1002250	1.06	2.09	5.2	3.95	6.82	6.01		
MAMMA1002253	2.77	2.39	3.45	4.84	6.18	3.37		
MAMMA1002267	17.17	19.95	51.7	130.02	108.53	115.75	**	+
MAMMA1002268	1.72	2.28	5.82	6.92	11.3	6.52		
MAMMA1002269	0.89	0.73	2.25	2.32	2.58	1.67		
MAMMA1002282	0.86	1.09	4.95	5.87	5.31	6.81		
MAMMA1002292	2.71	2.25	7.77	10.57	10.52	11.53	*	+
MAMMA1002293	3.71	3.31	12.81	8.54	10.47	12.05		
MAMMA1002294	0.9	1.71	4.61	3.68	6.03	4.2		
MAMMA1002297	1.53	3.25	7.45	5.77	7.8	6.91		
MAMMA1002298	1.48	1.4	3.98	3.85	3.11	2.46		
MAMMA1002299	1.5	1.69	3.16	3.91	2.97	2.2		
MAMMA1002308	1.39	1.35	6.55	4.5	3.11	2.54		
MAMMA1002310	3.56	3.84	12.73	9.92	12.66	11.48		

(1079)

特開2002-191363

2155

2156

MAMMA1002311	2.52	2.13	6.82	9.61	9.66	6.9	*	+
MAMMA1002312	1.63	2.22	5.19	3.51	8.45	2.55		
MAMMA1002317	2.08	2.55	4.89	4.08	3.85	4.09		
MAMMA1002319	0.8	2.78	3.51	2.68	3.97	2.85		
MAMMA1002322	2.48	3.23	7.84	12.21	10.02	8.55	*	+
MAMMA1002329	1.64	1.67	2.93	2.9	3.3	2.76		
MAMMA1002332	2.17	2.38	4.58	5.98	4.14	3.05		
MAMMA1002333	1.7	1.74	4.19	5.35	5.07	3.54		
MAMMA1002335	1.75	2.72	8.53	6.93	11.32	4.23		
MAMMA1002339	2.09	2.42	7.34	5.21	7.5	5.14		
MAMMA1002347	1.7	2.3	6.39	5.5	5.32	4.64		
MAMMA1002351	2.08	2.68	5.74	3.03	4.48	4.84		
MAMMA1002352	1.27	2.28	3.66	3.53	4.63	2.8		
MAMMA1002353	4.46	2.5	5.84	5.95	4.19	4		
MAMMA1002355	3.97	3.38	8.37	7.98	7.31	8.57		
MAMMA1002356	2.18	1.49	4.36	5.43	4.13	3.75		
MAMMA1002359	3.95	3.35	16.09	23.81	24.53	19	*	+
MAMMA1002360	0.93	1.73	3.77	2.48	3.2	1.67		
MAMMA1002361	2.01	2.64	4.53	4.17	4.95	4.03		
MAMMA1002362	2.33	2.33	3.36	5.31	5.51	3.99	*	+
MAMMA1002367	2.97	3.64	14.63	18.34	21.06	21.56	*	+
MAMMA1002371	2.28	3.75	8.3	6.15	6.74	5.88		
MAMMA1002380	1.81	2.26	4.9	4.71	5.76	3.55		
MAMMA1002384	2.14	1.53	4.73	4.48	5.36	4.05		
MAMMA1002385	1.19	2.05	5.63	3.34	4.8	2.47		
MAMMA1002390	1.41	2.04	3.75	5.48	5.4	3.43		
MAMMA1002392	1.94	3.1	6.1	4.06	5.95	3.32		
MAMMA1002396	4.87	3.49	12.87	10.79	12.9	8.08		
MAMMA1002399	4.42	5.13	10.69	10.95	8.66	4.57		
MAMMA1002400	3	2.22	4.69	3.11	4.36	3.53		
MAMMA1002409	51.57	55.16	63.3	77.54	80.62	77.88	**	+
MAMMA1002411	1.08	1.88	4.13	3.43	5.49	1.92		
MAMMA1002413	2.02	3.01	9.19	5.93	7.17	6.75		
MAMMA1002417	1.83	2.24	4.87	3.45	4.25	2.63		
MAMMA1002427	1.5	2.38	4.54	4.78	5.56	3.41		
MAMMA1002428	2.47	2.26	5.38	4.46	5.11	4.28		
MAMMA1002433	1.74	2.18	6.84	6.72	6.96	6.22		
MAMMA1002434	2.94	2.4	7.38	5.34	4.65	5.03		
MAMMA1002446	1.39	2.34	5.62	3.98	5.84	5.96		
MAMMA1002447	2.51	1.38	6.4	5.11	6.26	5.45		
MAMMA1002454	7.77	9.16	18.07	21.71	17.12	18.35		
MAMMA1002461	2.06	4.11	7.7	4.92	5.41	6.47		
MAMMA1002463	3.28	3.32	8.09	6.98	7.82	5.39		
MAMMA1002464	16.58	16.77	20.05	19.41	20.41	18.09		
MAMMA1002466	9.48	9.89	14.22	14.58	15.75	13.93		
MAMMA1002470	1.39	1.51	5.13	3.54	5.01	3.73		
MAMMA1002475	0.72	1.85	5.03	3.86	5.17	4.65		
MAMMA1002480	0.66	1.21	2.31	1.68	2.84	2.03		
MAMMA1002485	29.98	27.24	46.09	64.83	74.9	80.68	**	+
MAMMA1002494	2	2	4.11	4.48	5.12	5.13	*	+

(1080)

特開 2002-191363

2157

2158

MAMMA1002498	0.97	2.57	3.16	2.07	3.18	1.55		
MAMMA1002524	3.04	2.96	6.43	5.18	7.34	6.1		
MAMMA1002530	2.5	3.24	4.88	3.17	4.41	2.55		
MAMMA1002538	2.34	2.38	5.62	5.46	5.13	4.91		
MAMMA1002545	2.37	2.64	6.26	4.56	6.49	4.56		
MAMMA1002554	1.96	1.42	5.43	5.3	6.01	7.81		
MAMMA1002556	1.3	1.9	3.6	3.73	5.75	3.89		
MAMMA1002561	2.3	2.99	7.19	8.13	10.46	7.98		
MAMMA1002565	1.22	2.15	3.52	2.57	4.51	2.55		
MAMMA1002566	0.98	1.87	6.21	1.65	4.7	3.9		
MAMMA1002571	0.53	1.8	3.06	1.43	3.1	4.3		
MAMMA1002573	2.14	1.86	7.06	4.54	5.66	5.97		
MAMMA1002576	118.77	131.84	363.97	348.62	471.73	358.66		
MAMMA1002584	3.52	2.27	11.91	12.86	17.82	13.46		
MAMMA1002585	0.76	1.86	4.38	1.85	3.6	5.26		
MAMMA1002586	1.98	2.55	3.85	4.12	5.02	3.3		
MAMMA1002589	1.08	1.26	2.44	2.36	5.06	3.19		
MAMMA1002590	1.01	1.57	5.87	2.58	6.75	4.57		
MAMMA1002593	2.48	2.48	4.89	4.18	4.07	3.04		
MAMMA1002597	2.47	2.52	7.25	8.06	9.48	8.78	*	+
MAMMA1002598	12.12	13.52	30.83	37.28	48.14	38.91	*	+
MAMMA1002603	1.2	1.39	3.69	3.25	6.24	4.35		
MAMMA1002612	3.51	3.39	12.6	7.66	7.78	9.76		
MAMMA1002617	4.3	3.41	10.15	6.3	7.29	10.05		
MAMMA1002618	1.68	2.27	4.02	2.76	3.59	3.91		
MAMMA1002619	2.96	2.8	5.24	3.22	5.88	3.49		
MAMMA1002622	2.51	2.12	8.02	7.1	7.18	7.15		
MAMMA1002623	2.31	2.21	6.27	5.89	6.17	6.19		
MAMMA1002625	1.32	1.3	3.23	2.3	6.42	2.6		
MAMMA1002627	0.98	0.82	2.93	0.6	1.29	0.21		
MAMMA1002629	1.8	2.23	6.09	5.03	6.74	7.02		
MAMMA1002631	1	1.86	3.61	3.07	4.55	2.97		
MAMMA1002633	6.61	7.44	21.47	19.33	24.55	21.53		
MAMMA1002636	1.02	2.46	6.97	6.79	8.77	9.25		
MAMMA1002637	1.05	1.4	4.66	3.39	4.85	4.28		
MAMMA1002646	1.69	0.8	3.32	2.33	2.86	1.53		
MAMMA1002648	10.51	14.07	21.18	42.29	31.45	39.76	**	+
MAMMA1002650	1.33	0.56	1.62	1.76	2.08	0.57		
MAMMA1002652	1.76	2.82	7.31	7.5	7.41	9.79		
MAMMA1002655	1.7	2.11	3.65	2.54	4.23	3.78		
MAMMA1002662	0.84	2.24	4.33	3.57	5.68	4.13		
MAMMA1002665	3.61	3.57	10.05	13.42	17.97	19.59	*	+
MAMMA1002671	2.84	3.63	10.17	17.04	16.47	19.3	**	+
MAMMA1002673	1.32	2.14	4.93	4.07	5.03	2.82		
MAMMA1002684	2.95	3.11	3.84	6.61	8.19	7.54	**	+
MAMMA1002685	0.68	1.49	2.57	2.05	3.74	2.97		
MAMMA1002692	1.28	1.96	5.45	2.46	4.14	3.62		
MAMMA1002693	1.84	4.18	8	4.63	7.68	6.61		
MAMMA1002698	0.99	1.91	4.05	2.92	4.42	3.3		
MAMMA1002699	2	2.35	4.43	4.05	5.22	3.64		

(1081)

特開2002-191363

2159

2160

MAMMA1002701	2.41	2.56	8.46	6.72	8.94	8.93		
MAMMA1002708	1.51	1.55	5.38	4.08	6.16	6.18		
MAMMA1002711	1.58	2.08	7.04	4.37	7.35	5.81		
MAMMA1002712	3.05	3.13	6.98	4.88	7.12	7.39		
MAMMA1002716	0.56	1.75	3.39	2.38	6.29	2.9		
MAMMA1002721	2.11	2.01	5.57	3.72	6.34	4.59		
MAMMA1002723	2.43	2.46	4.91	3.85	5.98	4.88		
MAMMA1002727	3.85	5.55	5.78	5.29	4.45	6.22		
MAMMA1002728	21.35	22.03	57.81	49.09	54.73	65.13		
MAMMA1002742	4.12	4.39	10.35	7.92	8.63	7.61		
MAMMA1002743	4.12	3.89	6.17	13.81	14.09	13.46	**	+
MAMMA1002744	2.07	3.15	9.18	9.33	12.98	13.16		
MAMMA1002746	0.93	1.28	3.09	2.29	4.31	1.68		
MAMMA1002748	2.71	2.65	4.52	7.15	5.86	4.72	*	+
MAMMA1002754	1.12	2.41	5.56	5.05	5.65	6.26		
MAMMA1002758	0.71	1.66	2.55	1.57	4.41	1.69		
MAMMA1002762	11.3	11.14	36.64	38.42	34.23	48.71		
MAMMA1002764	1.83	3.2	5.95	5.11	6.06	4.26		
MAMMA1002765	1.19	1.63	4.29	4.63	5.26	2.67		
MAMMA1002769	7.4	6.44	13.04	13.78	8.03	12.41		
MAMMA1002771	1.41	2.41	3.31	3.54	5.39	4.39		
MAMMA1002775	4.56	4.48	19.79	22.54	29.77	24.29	*	+
MAMMA1002780	2.59	1.83	3.03	2.11	4.89	3.78		
MAMMA1002782	1.43	2.49	3.85	2.51	4.79	4.11		
MAMMA1002795	1.89	2.03	3.46	6.45	7.68	5.35	**	+
MAMMA1002796	4.35	3.97	7.51	7.2	8.09	8.17		
MAMMA1002805	6.61	11.12	16.52	15.95	24.7	16.5		
MAMMA1002806	1.47	2.02	3.51	2.28	4.62	2.17		
MAMMA1002807	1.63	2.4	6.77	6.78	9.66	6.4		
MAMMA1002814	3.43	3.52	7.92	9.58	12.39	10.66	*	+
MAMMA1002817	1.28	1.56	2.87	2.89	5.43	2.91		
MAMMA1002820	1.66	1.93	2.61	2.52	4.77	2.21		
MAMMA1002830	67.67	70.46	130.59	165.92	139.33	187.18	*	+
MAMMA1002833	4.16	2.88	9.4	8.22	10.68	10.58		
MAMMA1002835	0.77	1.87	4.03	1.73	3.97	2.79		
MAMMA1002838	1.85	2.66	5.31	2.91	4.44	3.93		
MAMMA1002842	1	3.83	3.84	3.32	4.63	5.15		
MAMMA1002843	1.72	2.92	2.33	4.09	4.81	3		
MAMMA1002844	3.05	3.64	6.52	5.26	7.3	4.09		
MAMMA1002845	1.25	1.57	2.45	3.59	3.55	4.67	*	+
MAMMA1002857	92.1	106.97	208.17	209.17	202.29	249.13		
MAMMA1002858	317.94	188.78	378.89	560.7	620.76	724.33	**	+
MAMMA1002863	2.17	2.83	6.91	3.51	5.12	3.96		
MAMMA1002868	2.73	3.7	6.26	6.35	9.53	10.25	*	+
MAMMA1002869	5.43	6.83	26.64	22.68	30.03	29.85		
MAMMA1002871	0.61	1.7	1.78	1.9	3.8	1.97		
MAMMA1002875	1.9	2.59	3.99	4.48	6.35	4.06		
MAMMA1002879	8.42	9.2	14.19	22.55	23.63	27.96	**	+
MAMMA1002880	1.23	2.02	2.12	1.48	5.42	2.03		
MAMMA1002881	1.21	1.43	1.84	3.01	5.43	2.46		

(1082)

特開2002-191363

2161

2162

MAMMA1002885	0.96	1.59	2.71	2.6	3.26	1.59		
MAMMA1002886	2.63	2.52	3.9	6.01	5.37	7.05	**	+
MAMMA1002887	1.28	1.83	2.78	2.98	5.14	4.32	*	+
MAMMA1002890	0.79	1.7	4.05	4.39	4.8	4.01		
MAMMA1002892	1.35	2.45	4.98	6.64	6.24	5.84	*	+
MAMMA1002893	4.52	3.58	5.4	7.6	8.03	8.43	**	+
MAMMA1002895	1.43	1.31	3.28	1.81	3.89	1.64		
MAMMA1002898	0.53	1.67	4.15	2.69	4.72	1.42		
MAMMA1002905	1.32	1.58	2.51	4.1	5.01	3.87	**	+
MAMMA1002906	15.12	10.76	15.42	19.47	13.76	15.58		
MAMMA1002908	0.99	1.24	4.28	3.53	4.24	4.07		
MAMMA1002909	1.92	2.64	5.67	6.82	8.18	6.57	*	+
MAMMA1002918	2.75	2.69	5.42	5.27	7.26	6.58		
MAMMA1002925	92.88	85.77	163.7	127.31	122.97	178.98		
MAMMA1002926	6.08	6.31	16.25	16.64	19.48	19.9		
MAMMA1002930	1.21	1.59	5.67	4.88	8.91	4.21		
MAMMA1002937	4.91	3.87	30.71	40.45	75.17	61.59	*	+
MAMMA1002938	1.67	1.86	2.42	2.35	3.2	3.56		
MAMMA1002941	0.49	1.48	2.78	2.53	3.59	2.24		
MAMMA1002947	2.24	2.59	4.55	6	6.8	7.94	*	+
MAMMA1002964	1.73	2.9	5.91	6.91	7.24	7.16	*	+
MAMMA1002967	1.94	1.59	2.28	2.9	4.19	2.79		
MAMMA1002970	2.72	1.77	6	7.59	7.28	8.96	*	+
MAMMA1002971	1.52	1.6	2.9	2.51	7.27	3.93		
MAMMA1002972	1	1.32	2.95	1.74	4.56	2.12		
MAMMA1002973	1.38	2.45	6.73	4.36	6.72	6.78		
MAMMA1002979	55.6	60.16	121.72	134.02	101.19	107.19		
MAMMA1002982	0.53	1.98	2.28	2.04	3.28	1.9		
MAMMA1002987	1.56	2.11	5.56	3.14	5.55	4.14		
MAMMA1003003	0.77	2.18	4.78	4.46	6.47	5.08		
MAMMA1003004	1.65	1.86	3.7	3.64	3.59	3.16		
MAMMA1003007	0.69	1.16	2.73	1.88	3.7	2.32		
MAMMA1003011	1.56	1.8	3.67	3.77	5.41	3.94		
MAMMA1003013	3.67	5.57	39.41	47.56	59.11	54.29	*	+
MAMMA1003015	1.16	1.8	2.21	2.54	2.9	2.19		
MAMMA1003019	0.6	1.61	2.1	3.12	4.61	2.63		
MAMMA1003020	2.96	4.19	5.34	11.31	10.33	10.09	**	+
MAMMA1003026	1.29	1.56	2.95	2.66	4.25	2.25		
MAMMA1003031	0.61	1.71	5.64	4.13	5.85	5.89		
MAMMA1003033	1.34	1.65	4.13	2.84	5.11	3.64		
MAMMA1003035	1.66	2.5	5.44	5.12	7.03	4.9		
MAMMA1003039	0.95	0.75	3.31	2.15	4.73	2.48		
MAMMA1003040	1.38	2.54	5.32	4.57	7.47	7.43		
MAMMA1003044	2.36	2.96	6.52	4.29	6.41	5.99		
MAMMA1003047	1.82	3.67	7.61	5.74	7.05	7.13		
MAMMA1003049	0.47	1.72	2.03	1.08	1.56	1.45		
MAMMA1003055	1.24	1.67	4.92	3.77	5.14	3.44		
MAMMA1003056	0.9	0.91	1.85	1.22	2.26	1.02		
MAMMA1003057	2.53	3.34	6.76	7.25	9.2	5.01		
MAMMA1003066	1.65	2.06	4.73	4.1	7.08	5.07		

(1083)

特開 2002-191363

2163

2164

MAMMA1003075	1.11	1.71	3.16	1.85	4.37	2.32		
MAMMA1003089	1.69	2.11	7.13	7.85	8.66	7.43		
MAMMA1003092	1.25	1.79	3.21	2.62	4.08	1.76		
MAMMA1003095	2.27	3.33	5.4	7.24	8.57	5.34		
MAMMA1003099	1.88	2.51	4.95	4.09	6.45	4.35		
MAMMA1003102	1.33	2.04	2.88	3.2	3.27	2.39		
MAMMA1003104	0.64	1.07	3.17	2.15	3.25	1.56		
MAMMA1003113	4.22	4.21	6.98	9.22	7.02	7.07		
MAMMA1003126	12.93	14.72	20.89	19.28	12	15.63		
MAMMA1003127	2.95	3.14	5.91	3.88	6.12	5.19		
MAMMA1003131	2.82	3.51	4.86	3.82	5.91	5.88		
MAMMA1003135	3.66	4.65	7.61	2.33	4.04	2.64		
MAMMA1003140	0.73	2.01	3.59	2.3	3.32	1.89		
MAMMA1003146	2.08	2.24	3.89	3.17	5.09	3.3		
MAMMA1003150	1.18	1.8	3.01	3.37	4.29	3.45	*	+
MAMMA1003154	0.54	1.41	2.29	2.21	3.57	2.34		
MAMMA1003155	8.08	9.18	20.88	21.41	17.68	20.56		
MAMMA1003157	5.94	4.82	6.07	5.18	7.05	7.89		
MAMMA1003163	1.74	1.69	4.23	2.55	5.46	3.08		
MAMMA1003164	2.94	4.56	6.23	4.08	9.9	8.18		
MAMMA1003166	3.62	3.5	5.77	6.24	8.66	6.12		
NB9N31000010	2.5	3.88	7.58	9.63	12.26	9.5	*	+
NB9N31000016	0.73	2.8	5.21	4.04	4.41	3.15		
NB9N31000043	8.1	8.88	19.71	12.51	12.3	12.64		
NB9N31000045	167.24	153.32	255.96	401.78	320.53	296.06	*	+
NB9N31000054	7.29	4.42	11.75	11.15	11.87	13.43		
NB9N31000076	2.31	1.94	3.51	4.5	6.35	4.44	*	+
NB9N31000086	2.62	2.65	6.23	3.61	9.71	7.69		
NT2RM1000001	2.56	2.45	6.24	5.7	7.05	6.32		
NT2RM1000018	3.84	4.69	10.6	6.58	9.09	6.92		
NT2RM1000032	1.12	2.64	3.88	2.28	4.92	3.21		
NT2RM1000035	1.72	3.68	5.53	5.44	5.21	5.98		
NT2RM1000037	1.38	2.98	2.75	2.41	4.15	2.11		
NT2RM1000039	3.45	5.13	5.9	6.51	7.26	8.4	*	+
NT2RM1000042	33.96	32.7	65.25	57.46	67.15	64.39		
NT2RM1000055	0.85	1.74	3.34	1.16	3.55	1.16		
NT2RM1000059	3.26	3.16	7.66	4.69	5.97	5.78		
NT2RM1000062	1.13	1.21	1.9	3.47	4.33	2.46	*	+
NT2RM1000065	23.8	16.41	34.06	36.15	35.1	51.38		
NT2RM1000066	4.13	4.31	8.98	7.23	10.95	9.81		
NT2RM1000071	49.63	37.81	86.71	73.04	63.32	84.05		
NT2RM1000080	1.37	2.04	3.8	5.1	5.94	4.5	*	+
NT2RM1000086	4.04	4.65	4.08	5.01	6.23	5.58	*	+
NT2RM1000092	6.17	6.93	15.76	14.48	25.91	15.13		
NT2RM1000118	0.63	1.12	1.22	0.63	1.7	0.44		
NT2RM1000119	1.32	2.27	1.96	1.84	3.38	2.99		
NT2RM1000121	1.13	1.84	1.76	2.92	3.84	2.78	*	+
NT2RM1000122	3.5	3.78	7.34	5.5	8.86	9.57		
NT2RM1000127	0.69	1.34	1.47	2.14	3.36	3.32	*	+
NT2RM1000131	0.71	1.7	1.47	1.36	3.02	2.53		

(1084)

特開2002-191363

2165

2166

NT2RM1000132	3.2	4.88	4.83	6.86	6.46	6.31	*	+
NT2RM1000153	1.75	1.9	3.68	2.38	4.45	4.84		
NT2RM1000184	72.82	77.46	151.91	106.39	163.07	125.55		
NT2RM1000186	1.55	1.46	4.32	2.67	4.72	3.94		
NT2RM1000187	3.11	1.96	5.16	10.09	9.1	8.78	**	+
NT2RM1000199	1.12	1.37	2.11	2.41	3.51	2.72	*	+
NT2RM1000213	1.32	1.75	2.38	2.66	2.71	2.22		
NT2RM1000215	10.95	11.07	17.21	19.51	22.84	15.14		
NT2RM1000218	9.72	9.95	23.71	26.94	24.74	29.21		
NT2RM1000224	8.8	8.63	15.2	15.51	21.29	17.61		
NT2RM1000236	30.38	24.19	61.14	72.86	82.44	71.5	*	+
NT2RM1000242	0.23	1.17	1.32	0.39	2.27	0.12		
NT2RM1000244	1.41	1.48	3.43	6.9	3.69	6.7	*	+
NT2RM1000252	1.75	1.5	3.4	3.4	3.06	3.18		
NT2RM1000256	7.88	5.89	9.46	26.12	29.45	36.8	**	+
NT2RM1000257	1.98	3.01	5.09	4.64	6.83	6.65		
NT2RM1000260	7.9	7.01	13.32	9.18	12.49	11.77		
NT2RM1000269	3.87	2.87	5.12	6.63	9.78	3.87		
NT2RM1000271	0.71	0.8	1.87	0.46	2.47	0.51		
NT2RM1000272	117.67	92.26	202.95	249.32	333.98	356.74	*	+
NT2RM1000273	10.03	9.45	20.12	22.32	16.68	15.76		
NT2RM1000274	63.11	66.41	123.01	137.14	91.97	104.48		
NT2RM1000280	3.95	4.18	8.18	6.71	8.72	7.93		
NT2RM1000295	0.49	1	2.2	1.12	3.16	0.87		
NT2RM1000300	1.51	1.87	2.78	3.63	5.75	3.09		
NT2RM1000304	58.38	98.72	161.87	187.58	185.55	204.78	*	+
NT2RM1000314	1.8	2.12	3.6	3.84	4.07	4.33		
NT2RM1000318	12.6	14.04	20.81	35.01	29.96	29.8	**	+
NT2RM1000335	2.76	2.57	4.34	6.29	5.41	4.09		
NT2RM1000341	0.46	1.27	1.95	1.41	2.33	0.99		
NT2RM1000350	3.04	3.47	5.52	7.32	5.63	6.44		
NT2RM1000354	0.55	1.31	1.31	5.43	7.2	5.72	**	+
NT2RM1000355	30.24	31.5	56.85	74.62	50.25	61.33		
NT2RM1000361	3.63	3.87	7.23	14.39	20.29	18.78	**	+
NT2RM1000365	0.58	1.08	1.71	1.27	1.82	0.52		
NT2RM1000372	14.99	19.56	30.06	42.71	46.67	45.44	**	+
NT2RM1000377	2.04	2.18	9.66	13.38	14.74	13.48	*	+
NT2RM1000388	0.35	1.57	3.01	2.2	3.8	2.42		
NT2RM1000394	0.45	1.31	1.87	1.43	2.72	0.69		
NT2RM1000399	0.53	1.57	3.25	1.98	3.2	1.81		
NT2RM1000407	1.13	1.52	2.17	1.02	2.7	1.51		
NT2RM1000421	0.84	0.57	2.78	1.06	1.77	1.13		
NT2RM1000422	20.65	23.31	54.69	87.5	82.91	79.47	*	+
NT2RM1000430	1.22	1.57	2.01	3.2	3.67	2.95	**	+
NT2RM1000462	1.55	2.33	7.32	5.59	7.28	8.16		
NT2RM1000499	1.36	2.09	4.74	5	6.16	6.37	*	+
NT2RM1000512	12.49	13.22	19.22	10.54	14.15	19.84		
NT2RM1000519	33.96	37.54	55.78	31.14	29.25	47.55		
NT2RM1000527	7.97	8.92	37.68	55.15	60.19	46.68	*	+
NT2RM1000539	3.45	3.59	12.93	15.52	17.01	18.1	*	+

(1085)

特開2002-191363

2167

2168

NT2RM1000542	0.85	1.05	2.99	1.17	2.35	1.02		
NT2RM1000553	3.7	2.42	22.32	42.83	42.96	34.5	*	+
NT2RM1000555	11.3	11.6	23.97	34.11	29.67	22.76		
NT2RM1000558	2.09	5.34	9.74	9.56	16.24	14.29		
NT2RM1000563	1.47	2.42	3.36	4.07	5.58	3.95	*	+
NT2RM1000566	0.88	1.57	3.5	3.62	6.01	2.79		
NT2RM1000570	96.92	77.32	137.63	167.35	105.47	174.1		
NT2RM1000571	13.21	11.87	22.51	43.87	40.18	28.45	*	+
NT2RM1000574	0.84	2.15	2.55	2.15	3.07	1.67		
NT2RM1000580	1.37	2.18	4.07	5.15	7.98	2.96		
NT2RM1000620	2.61	2.95	8.2	8.35	9.58	7.26		
NT2RM1000623	1.25	1.2	2.38	1.75	2.81	0.62		
NT2RM1000630	0.79	2.28	2.39	1.68	3.51	1.67		
NT2RM1000633	30.97	39.36	36.34	54.43	44.6	43.59	*	+
NT2RM1000634	1.91	4.16	8.12	2.56	7.05	5.57		
NT2RM1000642	3.85	5.37	8.13	8.21	8.56	8.52		
NT2RM1000647	41.3	39.09	62.11	57.72	68.29	62.69		
NT2RM1000648	2.49	2.65	4.61	6.14	5.63	4.51		
NT2RM1000650	2.46	3.05	7.6	5.4	6.07	6		
NT2RM1000661	4.48	5.7	15.82	15.48	13.45	13.18		
NT2RM1000666	1	1.77	1.99	1.37	2.8	0.71		
NT2RM1000669	3.51	2.76	4.67	3.63	5.42	3.28		
NT2RM1000672	2.23	3.95	7.81	3.98	8.47	7.22		
NT2RM1000681	99.53	86.09	118.7	105.41	90.59	124.14		
NT2RM1000691	2.02	2.61	5.74	3.61	7.69	3.76		
NT2RM1000698	1.11	1.43	4	6.42	6.29	4.11	*	+
NT2RM1000699	1.85	2.86	3.17	3.67	4.35	4.15	*	+
NT2RM1000702	3.71	4.64	9.47	9.31	9.72	11.4		
NT2RM1000703	11.56	12.36	25.24	26.72	20.42	21.06		
NT2RM1000704	24.48	23	32.91	46.54	24.13	40.82		
NT2RM1000725	60.92	59.45	88.28	94.89	82.36	105.67		
NT2RM1000726	1.85	2.02	5.75	1.97	4.8	4		
NT2RM1000731	1.11	2.24	4.98	2.45	3.47	3.43		
NT2RM1000741	1.38	1.87	3.16	2.69	4.15	2.9		
NT2RM1000742	2.61	4.6	7.41	9.55	10.94	9.84	*	+
NT2RM1000744	2.1	3.61	7.14	4.05	5.24	5.05		
NT2RM1000746	2.25	2.47	2.95	2.22	4.01	3.89		
NT2RM1000747	23.34	23.92	46.23	44.66	50.12	55.15		
NT2RM1000752	3.83	2.36	4.62	3.95	4.88	3.46		
NT2RM1000767	4.14	7.27	35.27	25.27	38.02	28.81		
NT2RM1000770	2.97	3.08	6.36	4.71	6.71	5.67		
NT2RM1000772	0.76	0.7	1.07	1.34	1.69	0.44		
NT2RM1000779	13.03	12.11	42.22	53.91	45.61	66.73	*	+
NT2RM1000780	1.16	2.9	3.74	3.09	4.32	3.01		
NT2RM1000781	1.07	0.98	1.71	2.58	4.4	1.93		
NT2RM1000789	5.28	5.15	29.74	29.63	46.72	36.53		
NT2RM1000800	2.87	2.63	6.37	5.66	9.57	6.96		
NT2RM1000802	2.44	2.99	7.5	4.34	5.47	4.82		
NT2RM1000811	1.78	1.6	2.13	2.26	4.96	2.76		
NT2RM1000826	6.06	6.36	13.34	14.42	20.73	20.98	*	+

(1086)

特開2002-191363

2169

2170

NT2RM1000829	3.91	2.87	6.39	6.73	8.48	8.41	*	+
NT2RM1000831	81.54	64.45	185.14	182.43	179.79	197.27		
NT2RM1000833	14.58	13.33	42.25	76.74	73.25	67.48	**	+
NT2RM1000834	4.06	3.09	6.2	8.49	9.42	10.49	**	+
NT2RM1000841	12.34	10.01	21.15	34.98	36.63	30.81	**	+
NT2RM1000848	4.79	4.42	6.44	9.36	12.74	10.45	**	+
NT2RM1000850	2.66	3.42	13.41	8.55	11.79	9.74		
NT2RM1000852	1.34	1.94	3.23	3.01	5.76	2.61		
NT2RM1000853	1.19	2.85	2.15	3.11	3.26	3.23		
NT2RM1000855	29.27	24.82	45.19	52.48	45.32	58.45		
NT2RM1000857	4.63	5	10.67	8.76	11.3	10.76		
NT2RM1000858	7.3	7.6	15.86	9.09	11.56	10.93		
NT2RM1000867	19.42	15.85	28.1	32.52	35.03	24.06		
NT2RM1000874	3.15	2.65	7.03	5.17	9.62	5.31		
NT2RM1000882	2.36	1.37	3.71	5.39	9.31	5.45	*	+
NT2RM1000883	5.21	3.34	7.42	5.18	11.3	7.9		
NT2RM1000885	3.86	4.43	9.4	7.59	8.15	9.8		
NT2RM1000893	3.15	3.41	8.14	7.73	6.17	8.39		
NT2RM1000894	3.29	4.4	6.18	8.14	6.11	6.61		
NT2RM1000898	3.72	7.33	10.02	13.4	17.51	12.41	*	+
NT2RM1000899	1.02	2.22	3.07	3.68	7.49	4.69		
NT2RM1000905	11.92	17.41	30.36	37.19	45.16	37.3	*	+
NT2RM1000910	7.5	8.78	20.16	36.37	36.98	37.5	**	+
NT2RM1000914	6.46	7.69	19.74	14.28	17.33	17.77		
NT2RM1000919	6.1	3.92	9.91	14.61	17.49	15.37	**	+
NT2RM1000921	0.72	1.9	3.69	2.79	4.27	3.32		
NT2RM1000922	4.7	6.11	8.09	9.03	5.21	6.36		
NT2RM1000924	0.89	3.03	3.04	3.08	2.89	3		
NT2RM1000927	1.35	1.78	2.85	3.07	4.72	3.46		
NT2RM1000951	7.95	11.33	26.73	32.33	34.46	31.18	*	+
NT2RM1000956	7.91	6.36	13.35	23.61	27.46	21.91	**	+
NT2RM1000960	12.48	10.27	29.06	34.95	37.47	38.96	*	+
NT2RM1000961	3.28	3.61	7.45	9.44	13.18	8.11		
NT2RM1000962	4.14	3.5	8.18	7.59	10.15	9.86		
NT2RM1000973	16.71	15.79	29.32	31.15	11.56	27.73		
NT2RM1000978	0.57	1.46	1.58	0.95	2.64	0.44		
NT2RM1000982	2.34	2.29	3.52	3.57	4.94	4.54	*	+
NT2RM1000991	1.61	1.78	4.25	3.88	5.56	5.23		
NT2RM1000994	6.36	6.16	12.57	16.52	16.64	14.53	*	+
NT2RM1001002	5.11	6.69	15.34	21.78	22.69	22.28	*	+
NT2RM1001003	5.42	5.15	11.98	16.24	9.06	8.46		
NT2RM1001008	1.4	2.22	2.48	1.83	4.34	4.33		
NT2RM1001011	6.29	5.43	7.86	14.4	10.46	14.72	*	+
NT2RM1001013	2.9	2.75	4.75	8.29	7.96	5.81	*	+
NT2RM1001017	1	1.82	3.44	3.28	4.86	3.92		
NT2RM1001018	65.15	74.45	146.86	134.65	125.46	113.93		
NT2RM1001026	1.37	2.64	3.17	2.99	4.61	3.31		
NT2RM1001028	0.98	1.73	2.91	1.74	1.89	0.76		
NT2RM1001043	4.47	3.64	8.42	11.43	12.7	8.01		
NT2RM1001044	2.23	3.17	4.92	5.03	5.51	3.93		

(1087)

特開2002-191363

2171

2172

NT2RM1001059	1.47	3.72	4.12	4.05	6.11	3.02		
NT2RM1001063	4.11	3.29	6.1	4.22	5.64	5.6		
NT2RM1001066	0.86	1.85	2.44	2.23	3.99	2.85		
NT2RM1001072	1.8	2.8	4.33	1.94	3.74	1.52		
NT2RM1001074	1.66	2.38	5.18	5.18	4.19	2.67		
NT2RM1001076	1.39	2.2	4.94	3.43	4.42	1.72		
NT2RM1001082	1.79	2.6	5.23	5.31	5.92	4.57		
NT2RM1001085	1.25	1.65	2.81	1.16	3.27	1.17		
NT2RM1001092	3.82	4.2	5.57	9.34	7.94	9.82	**	+
NT2RM1001102	1.7	2.3	4.4	2.49	5.94	4.64		
NT2RM1001103	4.37	3.88	7.18	6.25	10.28	8.08		
NT2RM1001105	1.77	2.02	4.63	2.49	5.11	3.51		
NT2RM1001112	2.68	2.66	3.69	3.85	4.75	2.43		
NT2RM1001115	1.44	1.57	4.72	3	6.46	3.73		
NT2RM1001122	2.84	3.35	7.3	9.43	9.75	9.54	*	+
NT2RM1001136	0.88	1.41	2.71	2.31	3.87	1.59		
NT2RM1001139	3.9	3.7	5.38	5.33	11.18	6.77		
NT2RM2000003	2.45	3.33	2.4	4.5	6.29	4.88	*	+
NT2RM2000006	2.34	2.95	7.25	5.12	7.11	6.24		
NT2RM2000010	12.79	13.03	22.58	20.2	17.11	21.83		
NT2RM2000013	8.1	9.44	50.36	68.75	95.32	74.36	*	+
NT2RM2000030	4.8	2.21	23.41	26.33	32.15	28.69		
NT2RM2000032	2.76	2.92	8.53	10.01	12.19	10.67	*	+
NT2RM2000039	3.94	4.67	4.75	6.42	5.78	4.99		
NT2RM2000042	3.5	4.9	11.69	17.71	17.4	15.02	*	+
NT2RM2000092	1	2.38	1.98	1.29	4.69	2.25		
NT2RM2000093	8.37	6.63	11.41	9.02	12.23	10.18		
NT2RM2000101	9.2	9.94	40	61.09	76.38	69.62	*	+
NT2RM2000104	6.82	8.02	46.75	51.34	68.83	43.48		
NT2RM2000124	1.54	2.23	6.33	7.73	8.84	8.47	*	+
NT2RM2000155	5.08	3.77	5.8	9.45	11.58	12.51	**	+
NT2RM2000191	3.33	5.68	28.62	26.54	34.38	31.6		
NT2RM2000192	1.03	1.29	2.45	6.3	4.75	3.83	*	+
NT2RM2000239	1.92	2.79	3.09	2.85	5.02	3.1		
NT2RM2000240	32.78	29.59	74.35	61.15	60.54	61.71		
NT2RM2000241	4.49	5.9	6.35	8.24	11.72	6.78		
NT2RM2000250	1.29	1.54	4.16	2.09	5.05	2.54		
NT2RM2000259	3.06	3.42	3.59	6.38	8.44	6.74	**	+
NT2RM2000260	2.53	2.05	3.12	4.23	4.07	5.79	*	+
NT2RM2000265	0.91	1.55	0.99	1.43	2.4	1.09		
NT2RM2000287	4.7	4.23	10.82	10.69	11.54	14.73		
NT2RM2000306	12.24	9.36	10.48	23.63	14	20.79	*	+
NT2RM2000312	19.4	17.81	25.01	38.39	31.27	24.8		
NT2RM2000322	1.93	1.82	4.48	3.79	7.05	3.32		
NT2RM2000343	7.74	8.38	41.34	63.81	79.6	71.12	*	+
NT2RM2000359	3.67	2.86	4.95	4.93	9.55	4.72		
NT2RM2000362	20.09	18.2	62.29	94.88	111.25	95.66	*	+
NT2RM2000363	1.08	1.89	2.97	4.2	4.32	3.33	*	+
NT2RM2000368	2.84	2.4	4.74	6.15	5.98	5.29	*	+
NT2RM2000371	76.64	65.68	119.32	135.82	125	44.64		

(1088)

特開2002-191363

2173

2174

NT2RM2000374	1.68	1.92	5.75	3.34	4.8	3.58		
NT2RM2000387	8.98	9.83	11.92	20.02	25.18	17.11	*	+
NT2RM2000393	1.7	1.63	3.75	3.31	7.65	3.28		
NT2RM2000395	1.07	1.51	1.98	1.72	4.34	2.23		
NT2RM2000402	12.38	11	15.78	25.15	18.31	22.51	*	+
NT2RM2000405	1.33	1.25	2.2	1.52	3.08	3.16		
NT2RM2000407	0.76	1.78	2.49	1.89	2.72	2.89		
NT2RM2000410	0.79	1.94	2.23	1.98	2.84	2.09		
NT2RM2000420	3.09	2.52	4.43	4.24	4.5	3.26		
NT2RM2000422	3.22	2.44	5.81	3.61	6.17	2.87		
NT2RM2000423	1.91	1.96	5.69	3.89	7.64	4.18		
NT2RM2000452	3.46	3.18	4.31	7.35	8.65	9.57	**	+
NT2RM2000469	3.28	3.28	4.44	1.87	2.33	2.46	*	-
NT2RM2000490	6.03	6.03	9.18	5.55	6.16	6.9		
NT2RM2000497	3.29	3.29	4.59	3.15	5.48	2.43		
NT2RM2000502	4.69	4.69	10.24	5.87	7.08	7.02		
NT2RM2000504	7.37	7.37	12.93	10.83	4.49	11.2		
NT2RM2000514	2.75	2.75	6.23	3.11	3.32	3.8		
NT2RM2000522	1.9	1.9	3.27	1.94	1.18	1.13		
NT2RM2000540	6.02	6.02	9.53	9.12	8.96	8.14		
NT2RM2000556	2.09	2.09	2.8	1.24	2.33	0.93		
NT2RM2000565	3.35	3.35	6.02	3.27	4.14	3.72		
NT2RM2000566	6.59	6.59	15.8	9.09	9.21	9.57		
NT2RM2000567	2.16	2.16	5.64	2	5.67	3.82		
NT2RM2000569	4.69	4.69	7.93	5.77	8.18	4.7		
NT2RM2000577	11.08	11.08	15.39	11.79	14.95	14.48		
NT2RM2000581	4.64	4.64	6.49	5.98	7.97	6.85		
NT2RM2000582	5.23	5.23	10.34	8.34	9.14	7.19		
NT2RM2000588	21.84	21.84	65.91	40.15	44.01	45.21		
NT2RM2000589	3.98	3.98	11.35	7.96	7.6	8.64		
NT2RM2000594	1.87	1.87	4.38	1.62	2.71	1.92		
NT2RM2000599	6.34	6.34	16.12	17.82	14.5	15.44		
NT2RM2000609	4.61	4.61	6.77	3.76	5.81	5.48		
NT2RM2000612	3.52	3.52	6.4	5.93	7.47	4.55		
NT2RM2000622	16.6	16.6	56.24	53.07	75.02	55.48		
NT2RM2000623	2.66	2.66	7.1	7.92	6.03	5.58		
NT2RM2000624	4.18	4.18	10.6	7.33	14.39	7.56		
NT2RM2000632	2.8	2.8	6.73	4.1	6.4	4.73		
NT2RM2000635	3.42	3.42	8.09	5.41	6.29	5.31		
NT2RM2000636	2.61	2.61	6.28	3.99	4.39	3.72		
NT2RM2000639	3.73	3.73	8.26	5.42	7.79	5.99		
NT2RM2000649	6.03	6.03	9.69	9.4	9.17	8.05		
NT2RM2000658	6.49	6.49	13.18	15.17	14.66	15.83	*	+
NT2RM2000660	11.45	11.45	18.34	17.03	7.1	20.16		
NT2RM2000669	3.6	3.6	6.51	5.28	4.28	6.69		
NT2RM2000689	31.07	31.07	59.7	37.03	16.51	70.9		
NT2RM2000691	2.09	2.09	5.73	4.83	7.13	4.27		
NT2RM2000714	3.41	3.41	10.97	11.46	14.54	11.3		
NT2RM2000718	4.08	4.08	7.15	2.88	5.42	4.33		
NT2RM2000732	5.38	5.38	14.81	9.49	14.18	8.25		

(1089)

特開2002-191363

2175

2176

NT2RM2000735	3.72	3.72	6.16	4.27	6.55	6.49		
NT2RM2000740	2.26	2.26	6.2	4.27	3.01	3.71		
NT2RM2000743	2.26	2.26	7.89	5.65	3.24	3.89		
NT2RM2000772	6.43	6.43	8.48	5.24	6.72	9.47		
NT2RM2000773	8.17	8.17	19.56	19.18	17.96	18.29		
NT2RM2000776	13.96	13.96	17.16	24.24	9.95	26.76		
NT2RM2000784	6.64	6.64	8.8	8.74	9.02	10.5		
NT2RM2000795	4.35	4.35	13.56	7.44	8.66	10.45		
NT2RM2000796	2.27	2.27	4.64	1.71	2.31	1.38		
NT2RM2000798	25.81	25.81	160.08	158.19	136.83	188.99		
NT2RM2000801	45.09	45.09	161.29	160.44	152.13	189.56		
NT2RM2000821	7.53	7.53	12.33	7.37	7.77	11.87		
NT2RM2000829	5.76	5.76	13.01	8.05	10.13	11.75		
NT2RM2000837	3.29	3.29	7.28	4.27	6.08	4.18		
NT2RM2000924	9.96	9.96	36.74	43.24	57.8	35.84		
NT2RM2000930	10.64	10.64	18.29	24.45	27.78	28.34	**	+
NT2RM2000937	4.35	4.35	8.62	5.08	6.66	6.56		
NT2RM2000939	1.12	1.12	2.37	2.67	1.84	1.82		
NT2RM2000942	124.8	124.8	253.61	161.4	118.61	210.11		
NT2RM2000951	1.01	1.01	2.6	2.13	3.12	1.6		
NT2RM2000952	2.53	2.53	5.31	5.98	6.41	6.86	*	+
NT2RM2000966	19.69	19.69	111.88	95.61	137.32	135.9		
NT2RM2000973	23.45	23.45	16.81	39.12	39.51	33.8	**	+
NT2RM2000983	10.07	10.07	18.59	30.68	39.13	27.52	*	+
NT2RM2000984	6.48	6.48	7.71	4.88	5.64	5.26	*	-
NT2RM2000994	8.27	8.27	16.47	13.2	8.36	21.81		
NT2RM2001004	6.01	6.01	48.58	47.51	54.15	46.8		
NT2RM2001022	101.09	101.09	350.92	239.63	304.37	490.04		
NT2RM2001035	10.75	10.75	24.98	24.17	24.54	34.51		
NT2RM2001038	5.77	5.77	9.86	10.48	11.75	7.9		
NT2RM2001043	4.45	4.45	10.02	5.1	7.41	6.74		
NT2RM2001050	2.71	2.71	6.89	4.72	5.83	4.03		
NT2RM2001055	3.78	3.78	5.89	4.24	6.31	4.7		
NT2RM2001065	6.17	6.17	15.91	8.51	12.12	14.21		
NT2RM2001075	39.81	39.81	188.21	154.65	156.72	168.68		
NT2RM2001083	2.23	2.23	5.57	4.01	5.8	3.01		
NT2RM2001100	10.38	10.38	93.67	95.8	113.9	97.52		
NT2RM2001105	6.34	6.34	8.27	11.35	5.34	11.8		
NT2RM2001109	6.81	6.81	9.4	11.88	12.47	14.53	*	+
NT2RM2001110	7.67	7.67	21.63	21.2	30.71	23.29		
NT2RM2001126	6.1	6.1	6.53	5.32	6.44	7.27		
NT2RM2001131	5.52	5.52	40.22	21.93	29.37	20.14		
NT2RM2001141	1.64	1.64	6.84	7.09	6.4	5.45		
NT2RM2001152	1.63	1.63	3.27	4.42	5.77	3.02		
NT2RM2001177	3.42	3.42	7.23	10.28	7.25	8.24		
NT2RM2001194	2.74	2.74	7.51	6.68	5.77	8.17		
NT2RM2001195	3.7	3.7	8.8	6.37	7.13	6.89		
NT2RM2001196	5.24	5.24	6.35	5.19	6.46	4.64		
NT2RM2001201	14.45	14.45	25.36	20.02	21.68	22.38		
NT2RM2001221	4.22	4.22	8.61	11.69	13.61	16.63	*	+

(1090)

特開2002-191363

2177

2178

NT2RM2001238	2.87	2.87	5.65	3.91	3.88	1.96		
NT2RM2001243	5.39	5.39	8.98	9.81	6.13	6.53		
NT2RM2001244	3.91	3.91	10.63	6.58	9.24	6.41		
NT2RM2001247	14.94	14.94	121.59	110.47	140.27	118.79		
NT2RM2001256	3.84	3.84	5.23	3.15	3.26	2.96		
NT2RM2001269	4.4	4.4	5.98	4.8	5.63	4.74		
NT2RM2001278	5.28	5.28	7.37	8.45	8.56	5.35		
NT2RM2001291	3.05	3.05	5.18	3.24	4.62	2.9		
NT2RM2001294	12.47	12.47	24.39	20.08	15.43	17.81		
NT2RM2001295	2.56	2.56	8.82	4.54	4.43	4.99		
NT2RM2001302	2.38	2.38	4.55	2.3	4.5	2.81		
NT2RM2001306	3.51	3.51	7.62	4.1	4.46	5.14		
NT2RM2001312	2.34	2.34	3.72	1.92	2.84	1.68		
NT2RM2001319	2.76	2.76	3.93	3.61	5.29	4.11		
NT2RM2001324	3.73	3.73	8.29	5.48	4.9	5.71		
NT2RM2001345	8.53	8.53	10.01	6.83	11.12	14.14		
NT2RM2001360	4.02	4.02	6.36	5.67	5.9	5.46		
NT2RM2001370	5.75	5.75	14.53	8.56	9.86	11.69		
NT2RM2001391	1.79	1.79	6.07	1.85	5.04	1.65		
NT2RM2001393	4.49	4.49	6.39	5.12	7.91	7.14		
NT2RM2001420	2.94	2.94	4.61	2.61	3.62	3.14		
NT2RM2001423	5.44	5.44	9.53	8.64	11.95	11.36		
NT2RM2001424	5.88	5.88	15.09	11.77	10.31	11.63		
NT2RM2001482	2.24	2.24	6.48	3.5	6.06	3.63		
NT2RM2001499	1.4	1.4	5.81	2.84	4.3	2.17		
NT2RM2001504	3.63	3.63	6.99	3.2	4.54	1.68		
NT2RM2001524	2.51	2.51	5.81	2.34	2.22	3.51		
NT2RM2001530	2.56	2.56	4.42	2.68	4.35	3.52		
NT2RM2001533	5.06	5.06	9.09	8.2	9.18	7.84		
NT2RM2001540	5.77	5.77	8.36	14.57	17.99	27.1	*	+
NT2RM2001544	2.4	2.4	6.12	3.7	3.72	2.31		
NT2RM2001547	6.6	6.6	15.29	8.44	7.61	8.24		
NT2RM2001558	1.53	1.53	3.44	1.76	4.87	1.71		
NT2RM2001575	2.45	2.45	4.57	3.36	4.38	2.29		
NT2RM2001582	2.99	2.99	4.98	2.2	5.16	3.06		
NT2RM2001588	3.69	3.69	8.8	6.39	9.14	6.6		
NT2RM2001592	2.66	2.66	6.2	3.1	5.24	4.64		
NT2RM2001603	4.74	4.74	8.7	10.42	12.03	11.77	*	+
NT2RM2001605	1.74	1.74	4.52	3.08	1.51	2.39		
NT2RM2001611	2.28	2.28	8.63	3.74	3.34	3.51		
NT2RM2001613	14.91	14.91	32.53	21.51	13.13	27.42		
NT2RM2001626	2.45	2.45	3.08	2.1	4.28	2.06		
NT2RM2001632	4.93	4.93	7.07	4.67	4.88	5.42		
NT2RM2001633	4.45	4.45	10.39	3.74	5.15	5.43		
NT2RM2001635	4.33	4.33	9.54	4.3	5.81	4.7		
NT2RM2001636	4.88	4.88	7.35	12.75	18.11	13.34	**	+
NT2RM2001637	1.25	1.25	6.48	4.18	3.68	2.51		
NT2RM2001639	3.98	3.98	9.32	4.67	4.33	3.29		
NT2RM2001641	1.63	1.63	4.69	4.84	6.02	2.71		
NT2RM2001643	2.78	2.78	7.46	4.79	4.4	2.83		

(1091)

特開2002-191363

2179

2180

NT2RM2001648	12.97	12.97	18.91	20.13	17.07	25.5		
NT2RM2001652	6.32	6.32	5.65	4.29	8.13	4.46		
NT2RM2001659	5.78	5.78	9.17	5.73	5.28	6.95		
NT2RM2001660	3.44	3.44	3.86	2.08	2.29	2.63	**	-
NT2RM2001664	1.24	1.24	6.12	4.51	4.89	4.8		
NT2RM2001668	3.72	3.72	8.16	7.66	5.72	7.02		
NT2RM2001670	1.62	1.62	4.11	2.88	3.96	3.56		
NT2RM2001671	2.67	2.67	5.57	3.9	6.46	4.85		
NT2RM2001675	1.94	1.94	4.28	1.97	3.73	0.64		
NT2RM2001681	2.47	2.47	5.91	3.13	4.64	3.39		
NT2RM2001685	4.58	4.58	5.68	1.29	2.72	1.14	**	-
NT2RM2001688	5.46	5.46	4.14	3.11	3.82	2.46	*	-
NT2RM2001695	15.09	15.09	35.18	17.41	19.26	34.51		
NT2RM2001696	2.74	2.74	6.64	7.15	6.7	6.8		
NT2RM2001698	1.44	1.44	3	4.06	3.49	1.65		
NT2RM2001699	1.63	1.63	5.03	4.19	3.75	5.48		
NT2RM2001700	1.65	1.65	4.13	2.56	3.37	3.91		
NT2RM2001704	2.68	2.68	5.46	3.89	3.85	3.99		
NT2RM2001706	4.29	4.29	6.77	3.33	3.13	3.32		
NT2RM2001714	6.48	6.48	6.64	5.62	7.33	5.18		
NT2RM2001716	0.97	0.97	3.7	3.03	5.49	2.92		
NT2RM2001718	1.91	1.91	3.47	5	3.5	3		
NT2RM2001723	2.09	2.09	5.48	5.1	5.21	5.71		
NT2RM2001727	3.08	3.08	6.25	7.51	7.28	5.7		
NT2RM2001730	3.52	3.52	7.15	5.04	5.43	3.85		
NT2RM2001738	4.56	4.56	6.2	6.71	10.25	9.08	*	+
NT2RM2001743	2.95	2.95	5.81	4.39	5.02	4.46		
NT2RM2001753	5.98	5.98	7.55	5.72	6.09	4.54		
NT2RM2001755	0.89	0.89	2.82	2.4	2.83	2.67		
NT2RM2001760	14.77	14.77	33.17	27.49	25.48	36.23		
NT2RM2001765	1.35	1.35	1.71	2.45	3.12	2.03	*	+
NT2RM2001767	12.04	12.04	120.66	148.84	168.4	146.29	*	+
NT2RM2001768	2.1	2.1	3.59	3.41	4.21	3.05		
NT2RM2001771	4.82	4.82	5.65	7.15	5.97	5.05		
NT2RM2001778	2.89	2.89	4.09	2.34	3.24	1.48		
NT2RM2001782	5.32	5.32	7.32	4.96	7.71	7.57		
NT2RM2001784	0.84	0.84	2.19	2.81	2.5	1.41		
NT2RM2001785	1.35	1.35	4.11	5.5	5.02	2.76		
NT2RM2001792	6.03	6.03	8.53	5.49	5.54	5.76		
NT2RM2001795	3.97	3.97	6.15	7.62	5.96	8.9		
NT2RM2001797	2.82	2.82	3.78	5	5.94	2.71		
NT2RM2001800	3.46	3.46	4.26	5.01	4.03	5.24		
NT2RM2001803	3.5	3.5	6.61	4.46	7.34	2.44		
NT2RM2001805	3.65	3.65	3.21	2.53	4.2	1.71		
NT2RM2001806	7.34	7.34	17.96	15.62	15.23	21.11		
NT2RM2001813	1.54	1.54	2.05	2.54	1.88	2.32		
NT2RM2001814	2.46	2.46	4.71	3.52	2.89	4.42		
NT2RM2001818	1.21	1.21	2.66	0.97	1.48	0.27		
NT2RM2001823	1.4	1.4	3.24	1.87	2.46	1.37		
NT2RM2001825	14.79	14.79	36.08	34.68	34.2	35.81		

(1092)

特開2002-191363

2181

2182

NT2RM2001832	5.93	5.93	6.1	5.19	5.48	2.93		
NT2RM2001839	67.48	67.48	123.46	152.63	157.11	121.35		
NT2RM2001840	3.04	3.04	7.13	4.61	5.11	5.37		
NT2RM2001851	3.92	3.92	7.61	3.78	4.74	6.49		
NT2RM2001855	8.21	8.21	11.51	10.22	12.06	15.41		
NT2RM2001867	2.82	2.82	5.01	2.83	5.62	3.74		
NT2RM2001869	60.8	60.8	90.58	101.19	79.67	105.32		
NT2RM2001879	3.01	3.01	6.99	2.55	3.19	2.66		
NT2RM2001883	1.52	1.52	3.26	0.98	2.28	0.73		
NT2RM2001886	1.57	1.57	4.56	2.48	3.51	2.42		
NT2RM2001887	3.78	3.78	7.66	4.48	4.97	5.73		
NT2RM2001896	274.2	274.2	378.57	325.68	216.52	497.31		
NT2RM2001902	1.92	1.92	4.28	1.31	2.92	2.14		
NT2RM2001903	16.25	16.25	42.55	35.47	31.71	37.22		
NT2RM2001930	2.11	2.11	6.3	2.3	5.86	5.44		
NT2RM2001935	4.16	4.16	5.04	3.16	4.42	5.87		
NT2RM2001936	2.81	2.81	4.9	2.99	3.44	4		
NT2RM2001939	3.56	3.56	3.34	1.82	3.01	3.16		
NT2RM2001941	1.84	1.84	4.29	2.84	2.82	2.72		
NT2RM2001950	4.66	4.66	10	6.01	6	8.69		
NT2RM2001952	2.67	2.67	4.78	2.49	4.55	5.37		
NT2RM2001976	11.48	11.48	18.2	14.58	11.46	35.27		
NT2RM2001982	1.85	1.85	3.91	2.04	2.47	1.88		
NT2RM2001983	4.45	4.45	8.36	4.18	6.49	7.54		
NT2RM2001984	7.74	7.74	8.88	20.06	27.05	22.95	**	+
NT2RM2001989	2.72	2.72	3.68	2.99	4.26	3.7		
NT2RM2001996	7.51	7.51	8.09	4.8	8.01	5.29		
NT2RM2001997	3.65	3.65	7.29	3.18	5.09	6.78		
NT2RM2001998	2.24	2.24	5.07	3.33	6.53	3.96		
NT2RM2001999	4.86	4.86	7.69	6.88	6.02	4.01		
NT2RM2002003	11.33	11.33	18.17	10.15	11	14.9		
NT2RM2002004	1.99	1.99	5.79	2.51	2.09	1.9		
NT2RM2002009	5.35	5.35	9.03	9.85	11.04	11.09	*	+
NT2RM2002014	2.62	2.62	3	3.65	4.47	4.03	**	+
NT2RM2002019	25.1	25.1	38.52	19.47	14.2	16.35		
NT2RM2002029	12.92	12.92	19.01	10.82	4.88	14.74		
NT2RM2002030	4.15	4.15	5.8	13.54	9.71	21.98	*	+
NT2RM2002034	22.05	22.05	31.76	24.83	20.89	21.04		
NT2RM2002049	7.4	7.4	12.12	9.76	10.42	13.22		
NT2RM2002055	2.8	2.8	8.01	1.91	4.03	2.52		
NT2RM2002072	9.26	9.26	12.88	12.28	19.12	12.82		
NT2RM2002088	4.82	4.82	13.85	11.35	11.7	15.28		
NT2RM2002091	4.98	4.98	8.44	5.92	3.82	4.61		
NT2RM2002100	3.26	3.26	6.05	4.82	4.24	3.19		
NT2RM2002109	1.31	1.31	3.57	2.57	4.88	4.92		
NT2RM2002126	21.41	21.41	32.24	35.28	22.31	31.52		
NT2RM2002128	3.7	3.7	5.17	2.74	3.86	2.41		
NT2RM2002129	6.43	6.43	11.48	8.53	13.03	10.66		
NT2RM2002142	5.72	5.72	9.74	5.26	8.91	6.4		
NT2RM2002144	3.27	3.27	3.76	1.85	1.73	1.62	**	-

(1093)

特開2002-191363

2183

2184

NT2RM2002145	2.63	2.63	8.69	6.1	5.18	5.98		
NT2RM2002153	2.61	2.61	6.37	6.31	7.62	5.75		
NT2RM2002163	0.97	0.97	3.41	1.87	3.4	0.64		
NT2RM2002170	3.28	3.28	7.03	6.62	7.5	7.65		
NT2RM2002178	3.99	3.99	3.67	3.5	5.48	2.9		
NT2RM2002179	7.82	7.82	8.69	6.17	8.15	6.02		
NT2RM2002270	4.51	4.51	4.56	2.28	1.76	1.67	**	-
NT2RM2002326	2.47	2.47	3.86	2.13	3.69	2.34		
NT2RM2002337	1.88	1.88	3.97	5.4	4.22	4.79	*	+
NT2RM2002339	2.83	2.83	6.29	5.26	5.22	3.85		
NT2RM2002345	5.16	5.16	6.03	4.04	4.2	4.21	*	-
NT2RM2002368	2.43	2.43	5.86	6.05	7.01	4.96		
NT2RM2002381	2.23	2.23	5.16	3.47	3.65	2.8		
NT2RM2002424	4.64	4.64	7.1	6.69	8.5	6.3		
NT2RM2002450	4.17	4.17	3.87	2.29	2.39	1.87	**	-
NT2RM2002482	3.93	3.93	4.65	2.66	3.2	3.79		
NT2RM2002492	9.39	9.39	24.31	29.13	24.65	29.29		
NT2RM2002575	3.26	3.26	5.23	5.99	6.03	5.07		
NT2RM2002580	4.23	4.23	4.68	4.82	7.79	7.42		
NT2RM2002592	7.7	7.7	12.59	13.07	15.28	14.69	*	+
NT2RM2002608	27.33	27.33	45.49	57.07	65.96	48.3	*	+
NT2RM2002615	6.01	6.01	9.38	13.15	20.32	14.42	*	+
NT2RM2002622	14.35	14.35	16.22	18.38	24.99	13.44		
NT2RM2002630	4.86	4.86	6.63	8.05	7.37	6.7		
NT2RM2002634	1.72	1.72	4.66	4.71	4.6	3.94		
NT2RM2002645	27.02	27.02	68.46	30.66	14.59	31.46		
NT2RM2002646	12.09	12.09	25.03	29.45	22.88	34.8		
NT2RM2002647	7.68	7.68	17.56	19.5	17.74	23.44		
NT2RM2002652	5.11	5.11	4.71	4.3	6.43	3.73		
NT2RM2002692	4.59	4.59	4.48	2.5	2.86	2.02	**	-
NT2RM2002721	30.26	30.26	46.01	62.71	84.18	61.02	*	+
NT2RM2002748	18.37	18.37	43.62	87.35	119.27	102.35	**	+
NT2RM2002764	2.28	2.28	5.3	5.05	5.95	4.07		
NT2RM2002772	3.15	3.15	9.32	9.66	7.81	5.44		
NT2RM2002811	5.79	5.79	12.3	14.01	9.18	10.45		
NT2RM2002818	2.03	2.03	7.94	5.86	5.42	7.13		
NT2RM2002879	4.21	4.21	7.17	8.39	7.87	9.11	*	+
NT2RM2002979	11.79	11.79	19.66	24.49	23.23	21.79	*	+
NT2RM2002981	4.42	4.42	3.78	3.58	4.95	2.63		
NT2RM2002995	5.13	5.13	3.29	3.42	3.74	3.5		
NT2RM2003031	1.37	1.37	2.63	3	2.58	1.9		
NT2RM2003042	4.1	4.1	10.77	10.59	6.02	6.07		
NT2RM2003044	1.88	1.88	4.11	2.13	5.11	1.2		
NT2RM2003090	4.4	4.4	7.64	9.36	7.91	10.68	*	+
NT2RM2003095	11.98	11.98	25.25	15.63	16.43	19.04		
NT2RM2003116	11.16	11.16	16.09	17.96	21.43	22.08	*	+
NT2RM2003222	3.98	3.98	3.63	2.67	3.64	2.35		
NT2RM2003224	11.29	11.29	15.33	24.29	29.77	20.76	*	+
NT2RM2003250	14.18	14.18	86.06	85.79	96.6	94.15		
NT2RM2003258	4.59	4.59	6.32	6.54	5.11	5.69		

(1094)

特開2002-191363

2185

2186

NT2RM2003262	5.07	5.07	7.33	5.06	7.76	5.72		
NT2RM4000023	2.15	2.15	7.02	3.57	4.91	4.29		
NT2RM4000024	2.28	2.28	6.78	3.17	4.98	4.33		
NT2RM4000027	4.74	4.74	7.77	4.85	6.94	11.32		
NT2RM4000030	2.95	2.95	5.73	3.16	4.9	2.64		
NT2RM4000033	2.51	2.51	4.77	2.36	3.89	4.61		
NT2RM4000034	1.93	1.93	5.35	3.74	4.84	6.09		
NT2RM4000046	1.37	1.37	3.79	1.57	3.5	2.32		
NT2RM4000052	1.82	1.82	3.55	1.72	2.96	1.98		
NT2RM4000054	10.43	10.43	13.85	12.07	12.83	24.7		
NT2RM4000061	1.65	1.65	4.17	1.66	4.54	0.83		
NT2RM4000074	15.83	15.83	43.57	27.9	34.24	30.79		
NT2RM4000085	5.35	5.35	10.1	8.41	10.19	10.17		
NT2RM4000086	3.06	3.06	4.5	3.84	5.25	3.71		
NT2RM4000100	6.62	6.62	15.05	12.74	15.6	14.84		
NT2RM4000101	3.77	3.77	9.11	7.17	7.71	8.78		
NT2RM4000102	32.35	32.35	42.47	27.24	19.34	70.54		
NT2RM4000104	2.78	2.78	7.13	3.41	5.12	4.3		
NT2RM4000115	2.87	2.87	6.1	3.91	5.86	4.08		
NT2RM4000129	2.17	2.17	4.75	2.62	3.48	2.18		
NT2RM4000139	3.17	3.17	3.31	3.58	4.82	6.49		
NT2RM4000149	2.74	2.74	1.49	2.41	2.55	7.32		
NT2RM4000155	2.73	2.73	5.5	2.13	4.51	3.71		
NT2RM4000156	5.94	5.94	16.74	20.45	21	21.96	*	+
NT2RM4000167	1.36	1.36	2.58	2.61	4.34	1.75		
NT2RM4000169	9.95	9.95	36.53	29.12	24.89	23.57		
NT2RM4000191	4.29	4.29	7.56	5.49	5.57	5.66		
NT2RM4000197	2.73	2.73	4.78	1.83	3.86	2.2		
NT2RM4000198	3.38	3.38	7.42	5.26	5.45	4.21		
NT2RM4000199	2	2	3.51	2.8	4	3.76		
NT2RM4000200	0.67	0.67	3.19	2.25	1.84	1.2		
NT2RM4000202	1	1	3.24	2.11	2.42	1.84		
NT2RM4000210	1.46	1.46	3.72	2.41	3.08	2.21		
NT2RM4000215	2.54	2.54	5.43	3.3	4.09	3.06		
NT2RM4000220	6.42	6.42	10.52	8.68	11.08	15.14		
NT2RM4000229	3.26	3.26	6.62	1.46	2.34	1.76		
NT2RM4000231	6.37	6.37	7.06	6.13	7.85	6.24		
NT2RM4000233	4.83	4.83	17.3	11.9	14.34	13.4		
NT2RM4000244	2.35	2.35	5.22	3.86	4.14	5.17		
NT2RM4000251	3.85	3.85	10.97	4.82	6.52	4.59		
NT2RM4000255	2.28	2.28	4.7	3.79	4.89	1.42		
NT2RM4000265	2.23	2.23	5.69	4.29	8.21	1.99		
NT2RM4000283	18.14	18.14	26.21	37.17	39.88	44.79	**	+
NT2RM4000284	13.85	13.85	33.72	31.96	42.88	40.67		
NT2RM4000290	6.31	6.31	7.76	3.77	4.92	4	*	-
NT2RM4000295	2.36	2.36	2.16	2.32	2.13	0.88		
NT2RM4000306	3.79	3.79	7.76	7.1	6.14	5.02		
NT2RM4000307	5.04	5.04	9.13	9.95	9.99	11.72	*	+
NT2RM4000309	2.48	2.48	5.34	3.92	6.52	5.52		
NT2RM4000313	3.92	3.92	9.61	5.75	7.77	8.52		

(1095)

特開 2002-191363

2187

2188

NT2RM4000318	3.38	3.38	6.87	4.35	6.36	3.28		
NT2RM4000324	4.93	4.93	5.93	2.79	4.98	2.12		
NT2RM4000326	5.32	5.32	4.61	2.59	2.45	2.01	**	-
NT2RM4000327	4.97	4.97	10.95	7.94	10.32	7.71		
NT2RM4000344	5.46	5.46	16.67	11.16	10.17	19.18		
NT2RM4000349	3.68	3.68	9.99	11.87	10.88	13.8	*	+
NT2RM4000354	1.65	1.65	3.13	4.2	4.31	3.1		
NT2RM4000356	1.5	1.5	3.11	2.5	4.07	1.64		
NT2RM4000366	15.75	15.75	44.48	38.81	44.07	58.06		
NT2RM4000368	3.04	3.04	5.9	4.36	5.48	3.48		
NT2RM4000373	6.49	6.49	12.29	12.72	15.96	16.47	*	+
NT2RM4000386	4.92	4.92	4.71	3.81	4.57	4.6		
NT2RM4000395	2.7	2.7	4.69	6.36	6.51	5.68	*	+
NT2RM4000414	1	1	2.76	2.38	2.9	2.19		
NT2RM4000417	1.66	1.66	2.83	3.9	3.95	3.25	*	+
NT2RM4000421	2.99	2.99	5.17	4.96	5.47	4.13		
NT2RM4000425	10.56	10.56	26.8	26.49	31.48	45.28		
NT2RM4000433	2.78	2.78	5.39	1.67	2.21	1.79		
NT2RM4000436	3.8	3.8	9.47	11.84	16.75	16.38	*	+
NT2RM4000444	4.51	4.51	12.97	7.29	8.54	7.38		
NT2RM4000457	3.35	3.35	8.69	13.35	12.38	13.25	*	+
NT2RM4000471	1.73	1.73	4.01	4.17	4.87	2.49		
NT2RM4000472	2.2	2.2	7.62	6.64	7.61	5.39		
NT2RM4000486	2.98	2.98	5.92	6.85	7.54	6		
NT2RM4000490	3.85	3.85	6.41	7.16	5.1	5.86		
NT2RM4000496	3.68	3.68	3.86	2.16	2.36	2.04	**	-
NT2RM4000505	26.85	26.85	60.33	68.9	80.59	70.67	*	+
NT2RM4000511	22.8	22.8	45.35	64.6	89.95	75.97	*	+
NT2RM4000514	2.61	2.61	6.75	10.47	7.53	9.25	*	+
NT2RM4000515	3.75	3.75	8.81	7.27	7.94	4.66		
NT2RM4000517	34.51	34.51	74.2	76.45	53.07	79.47		
NT2RM4000520	2.24	2.24	3.08	3.21	3.3	5.49		
NT2RM4000531	2.76	2.76	5.71	4.41	5.5	4.22		
NT2RM4000532	3.03	3.03	5.56	2.72	3.54	2.39		
NT2RM4000533	3.55	3.55	5.43	4.73	6.98	3.23		
NT2RM4000534	5.17	5.17	2.92	2.73	4.62	1.94		
NT2RM4000563	5.21	5.21	16.69	17.05	16.8	19.36		
NT2RM4000566	1.79	1.79	4.6	7.14	5.27	7.91	*	+
NT2RM4000568	10.48	10.48	19.4	28.18	22.91	33.06	*	+
NT2RM4000585	1.97	1.97	2.52	0.82	1.99	1.26		
NT2RM4000587	3.88	3.88	6.15	6.66	5.75	5.15		
NT2RM4000590	3.73	3.73	4.12	1.64	2.48	2.3	**	-
NT2RM4000593	4.46	4.46	7.83	10.21	12.36	7.45		
NT2RM4000595	3.94	3.94	3.91	2.27	4.36	2.45		
NT2RM4000603	4.18	4.18	6.98	6.95	7.23	7.26		
NT2RM4000611	9.81	9.81	18.02	17.1	17.38	25.49		
NT2RM4000616	2.05	2.05	6.19	2.73	4.14	4.53		
NT2RM4000621	26.04	26.04	70.86	57.5	62.2	59.07		
NT2RM4000648	2.78	2.78	7.83	2.66	2.67	3.88		
NT2RM4000649	5.53	5.53	11.03	7.34	8.4	13.12		

(1096)

特開2002-191363

2189

2190

NT2RM4000658	3.22	3.22	8.18	4.22	6.14	8.14		
NT2RM4000661	57.68	57.68	140.68	141.09	135.49	177.49		
NT2RM4000673	3.02	3.02	5.72	3.69	3.8	4.59		
NT2RM4000674	2.23	2.23	4.27	2.89	2.63	2.99		
NT2RM4000689	3.81	3.81	6.47	5.34	6.36	7.74		
NT2RM4000698	14.85	14.85	20.92	25.12	23.66	26.39	*	+
NT2RM4000700	2.39	2.39	5.53	2.41	6.95	5.61		
NT2RM4000701	10.07	10.07	54.36	61.81	67.75	63.17		
NT2RM4000712	3.5	3.5	7.9	5.97	8.03	9.55		
NT2RM4000717	2.14	2.14	6.66	3.67	2.94	4.3		
NT2RM4000733	4.37	4.37	7.8	4.16	6.93	11.03		
NT2RM4000734	2.17	2.17	5.92	2.35	5.23	4.7		
NT2RM4000741	2.14	2.14	6.11	3.59	4.75	4.66		
NT2RM4000744	1.76	1.76	7.05	2.76	4.4	10.18		
NT2RM4000749	15.53	15.53	23.13	26.26	27.8	34.67	*	+
NT2RM4000751	2.88	2.88	6.54	6.23	6.11	5.94		
NT2RM4000752	4.11	4.11	4.88	4.78	5.12	38.58		
NT2RM4000760	3.5	3.5	9.69	4.54	6.31	5.26		
NT2RM4000761	237.9	237.9	478.3	219.65	302.54	336.34		
NT2RM4000764	66.05	66.05	178	212.33	205.98	232.75	*	+
NT2RM4000768	6.11	6.11	11.21	15.56	10.14	21.17		
NT2RM4000778	1.6	1.6	4.7	4.27	5.18	6.18		
NT2RM4000779	4.52	4.52	8.28	6.87	7.19	7.33		
NT2RM4000787	2.55	2.55	7.49	3.64	4.9	4.53		
NT2RM4000790	2.99	2.99	5.03	5.47	5.82	12.06		
NT2RM4000795	1.99	1.99	3.67	2.36	1.2	2.51		
NT2RM4000796	3.26	3.26	5.86	4.29	3.48	4.28		
NT2RM4000798	1.77	1.77	5.53	3.72	3.08	3.47		
NT2RM4000800	4.15	4.15	8.16	8.7	9.44	9.06		
NT2RM4000813	3.31	3.31	8.79	7.14	7.95	10.09		
NT2RM4000820	4.89	4.89	9.14	5.39	6.27	5.44		
NT2RM4000827	7.1	7.1	18.55	16.3	15.8	17.88		
NT2RM4000830	3.27	3.27	7.35	5.28	7.8	8.38		
NT2RM4000833	2.51	2.51	6.84	4.48	3.35	4.87		
NT2RM4000841	4	4	15	10.57	8.84	10.1		
NT2RM4000846	1.66	1.66	8.83	4.74	6.83	5.09		
NT2RM4000848	2.61	2.61	5.75	4.15	6.32	3.12		
NT2RM4000852	3.89	3.89	9.81	8.16	8.67	8.29		
NT2RM4000855	5.12	5.12	7.64	5.99	6.08	7.19		
NT2RM4000859	11.18	11.18	16.28	16.48	19.12	17.62		
NT2RM4000868	3.06	3.06	6.47	5.23	6.91	7.05		
NT2RM4000870	4	4	9.82	7.06	3.87	6.59		
NT2RM4000879	1.67	1.67	6	4.15	3.31	3.11		
NT2RM4000882	18.99	18.99	28.36	20.33	16.5	12.55		
NT2RM4000887	2.16	2.16	6.01	3.53	4.91	1.6		
NT2RM4000895	2.33	2.33	5.33	3.16	5.9	3.08		
NT2RM4000897	5.78	5.78	6.99	8.89	9.73	7.02		
NT2RM4000901	5.22	5.22	6.41	4.37	6.2	6.12		
NT2RM4000950	4.04	4.04	5.09	3.57	4.06	2.91		
NT2RM4000965	2.89	2.89	4.54	5.98	4.04	6.18		

(1097)

特開 2002-191363

2191

2192

NT2RM4000971	2.49	2.49	4.78	4.84	4.29	7.62		
NT2RM4000979	5.16	5.16	12.71	10.23	10.46	7.87		
NT2RM4000987	1.9	1.9	4.59	3.64	4.22	3.21		
NT2RM4000989	2.51	2.51	5.17	4.2	5.16	4.99		
NT2RM4000991	3.1	3.1	4.83	2.87	3.53	8.73		
NT2RM4000992	3.39	3.39	5.41	3.6	4.54	3.19		
NT2RM4000996	6.22	6.22	7.86	7.14	8	7.28		
NT2RM4000997	3.53	3.53	13.96	9.81	10.96	10.89		
NT2RM4001001	26.06	26.06	55.43	37.67	34.22	54.29		
NT2RM4001002	5.13	5.13	11.03	11.54	11.33	19.44		
NT2RM4001016	1.63	1.63	2.73	4.07	5.31	3.76	*	+
NT2RM4001025	65.77	65.77	133.97	148.39	181.87	171.5	*	+
NT2RM4001027	2.49	2.49	3.66	1.67	1.77	4.31		
NT2RM4001032	2.55	2.55	6.74	4.94	5.46	3.84		
NT2RM4001047	3.87	3.87	3.7	2.61	2.73	2.7	**	-
NT2RM4001049	3.97	3.97	10.12	18.29	20.63	26.25	**	+
NT2RM4001051	2.72	2.72	12.54	11.17	10.12	13.68		
NT2RM4001052	14.95	14.95	72.14	75.49	75.01	79.12		
NT2RM4001053	14.96	14.96	39.3	41.36	28.95	25.87		
NT2RM4001054	3.13	3.13	5.17	5.34	5.69	5.01		
NT2RM4001059	3.65	3.65	6.37	4.91	3.52	4.48		
NT2RM4001071	4.03	4.03	7.35	6.8	7.34	6.55		
NT2RM4001084	8.04	8.04	6.52	9.49	10.53	9.53	*	+
NT2RM4001092	12.61	12.61	109.97	76.93	98.78	73.14		
NT2RM4001100	6.72	6.72	20.93	22.35	15.18	18.98		
NT2RM4001116	1.17	1.17	2.5	2.77	2.96	2.16		
NT2RM4001119	1.74	1.74	4.82	4.72	4.22	4.15		
NT2RM4001140	2.65	2.65	7.49	8.27	7.78	9.03		
NT2RM4001148	6.59	6.59	13.68	12.1	12.92	15.83		
NT2RM4001151	3.7	3.7	4.31	1.94	3.65	1.68		
NT2RM4001155	5.51	5.51	6.36	3.55	5.37	4.63		
NT2RM4001157	1.68	1.68	4.95	4.86	2.81	3.82		
NT2RM4001160	1.57	1.57	3.39	2.99	1.9	2.41		
NT2RM4001163	42.35	42.35	86.27	106.63	52.13	98.56		
NT2RM4001187	2.64	2.64	7.25	3.96	4.87	5.2		
NT2RM4001191	3.2	3.2	9.95	8.59	6.36	8.72		
NT2RM4001200	3.83	3.83	6.35	5.52	4.41	4.11		
NT2RM4001203	9.93	9.93	20.29	22.65	25.36	21.82		
NT2RM4001204	3.23	3.23	3.59	3.17	2.83	2.6		
NT2RM4001217	4.62	4.62	12.16	11.26	13.39	13.72		
NT2RM4001245	7.31	7.31	17.14	13.76	14.49	17.16		
NT2RM4001247	3.23	3.23	9.1	5.73	6.57	5.03		
NT2RM4001256	2.51	2.51	6.39	3.57	5.48	4.14		
NT2RM4001258	8.2	8.2	24.68	25.02	24.16	22.89		
NT2RM4001267	3.43	3.43	4.83	3.93	4.3	8.08		
NT2RM4001273	4.23	4.23	8.38	7.39	6.96	8.77		
NT2RM4001281	4	4	10.54	10.15	9.05	10		
NT2RM4001286	345.27	345.27	526.77	215.2	220.51	552.53		
NT2RM4001290	23.51	23.51	61.5	56.51	48.6	59.58		
NT2RM4001309	2.64	2.64	6.81	3.17	5.42	4.05		

(1098)

特開2002-191363

2193

2194

NT2RM4001313	2.63	2.63	8.62	3.81	6.36	7.03		
NT2RM4001316	3.14	3.14	6.12	3.39	3.85	5.25		
NT2RM4001320	2.4	2.4	6.43	2.83	3.16	5.6		
NT2RM4001321	3.98	3.98	8.62	6.17	7.03	6.24		
NT2RM4001325	2.54	2.54	5.2	4.76	2.93	5.25		
NT2RM4001333	8.65	8.65	18.06	8.57	10.76	9.51		
NT2RM4001340	4.81	4.81	12.27	6.99	7.72	11.24		
NT2RM4001344	4.09	4.09	4.69	3.04	3.44	4.93		
NT2RM4001347	6.49	6.49	9.8	10.53	9.72	17.71		
NT2RM4001357	7.59	7.59	12.09	8.58	11.68	9.37		
NT2RM4001360	2.79	2.79	5.11	3.82	2.85	3.07		
NT2RM4001371	4.71	4.71	8.57	6.58	13.16	12.93		
NT2RM4001377	8.01	8.01	13.74	8.66	9.23	10.58		
NT2RM4001382	17.31	17.31	56.74	35.57	28.32	38.22		
NT2RM4001384	2.17	2.17	4.11	2.35	3.34	3.75		
NT2RM4001400	3.78	3.78	5.76	4.87	4.66	5.07		
NT2RM4001409	2.55	2.55	6.82	4.82	4.63	5.27		
NT2RM4001410	5.48	5.48	21.69	17.34	21.09	18.91		
NT2RM4001411	2.66	2.66	6.5	5.86	6.64	6.77		
NT2RM4001412	1.75	1.75	3.33	2.39	2.52	2.79		
NT2RM4001414	2.18	2.18	4.83	2.45	2.4	3.77		
NT2RM4001436	8.35	8.35	16.65	10.45	10.13	15.55		
NT2RM4001437	2.77	2.77	8.85	8.31	11.26	7.67		
NT2RM4001444	11.57	11.57	25.93	21.1	18.36	18.42		
NT2RM4001454	3.62	3.62	7.89	6.16	5.03	4.79		
NT2RM4001455	5.85	5.85	22.19	18.03	25.08	26.16		
NT2RM4001483	4.37	4.37	8.77	4.97	5.43	6.46		
NT2RM4001489	3.12	3.12	7.04	3.64	4.19	5.46		
NT2RM4001495	1.35	1.35	5.13	2.85	3.29	3.37		
NT2RM4001499	1.74	1.74	4.98	3.65	2.77	3.3		
NT2RM4001515	0.95	0.95	3.38	2.65	5.38	3.41		
NT2RM4001519	1.86	1.86	4.6	2.24	4.38	2.85		
NT2RM4001522	3.8	3.8	7.66	5.36	7.42	9.13		
NT2RM4001523	5.46	5.46	9.11	4.82	7.58	5.79		
NT2RM4001550	11.9	11.9	17.38	16.49	16.76	15.01		
NT2RM4001553	7.88	7.88	13.4	23.26	23.53	23.87	**	+
NT2RM4001554	0.86	0.86	1.74	1.85	1.11	2.2		
NT2RM4001557	2.5	2.5	6.33	5.05	3.36	4.89		
NT2RM4001565	1.87	1.87	4.05	2.46	3.34	3.23		
NT2RM4001566	3.23	3.23	8.57	8.91	10.49	11.42		
NT2RM4001569	1.47	1.47	5.4	3.35	4.15	1.56		
NT2RM4001579	6.57	6.57	16.69	19.23	23.83	18.22		
NT2RM4001582	4.06	4.06	5.97	2.16	3.17	2.67	*	-
NT2RM4001589	21.51	21.51	37.16	42.45	55.76	47.57	*	+
NT2RM4001592	1.37	1.37	2.96	3.02	2.14	3.71		
NT2RM4001594	1.98	1.98	4.09	5.4	5.24	5.67	*	+
NT2RM4001597	2.65	2.65	5.64	5.17	4.97	4.33		
NT2RM4001605	2.7	2.7	6.18	5.71	5.92	5.93		
NT2RM4001609	23.65	23.65	45	61.08	78.89	77.31	*	+
NT2RM4001610	48.1	48.1	69.16	132.54	132.39	115.22	**	+

(1099)

特開2002-191363

2195

2196

NT2RM4001611	3.31	3.31	4.56	2.33	2.32	2.02	*	-
NT2RM4001618	7.05	7.05	7.95	6.68	8.98	12.95		
NT2RM4001622	13.53	13.53	19.88	14.67	24.67	28.46		
NT2RM4001624	1.6	1.6	3.02	3.92	2.66	4.43		
NT2RM4001625	4.89	4.89	39.6	41.63	47.1	46.46		
NT2RM4001629	3.82	3.82	8.82	12.09	12.08	13.38	*	+
NT2RM4001632	15.28	15.28	24.55	31.07	26.16	25.6		
NT2RM4001642	3.29	3.29	4.17	2.89	3.62	2.1		
NT2RM4001647	4.44	4.44	6.83	4.04	5.48	4.67		
NT2RM4001650	4.96	4.96	4.94	2.66	2.87	3.79	**	-
NT2RM4001662	2.18	2.18	5.47	8.31	6.54	9.39	*	+
NT2RM4001666	2.28	2.28	6.5	6.24	6.17	8.14		
NT2RM4001670	3.52	3.52	10.77	11.16	10.82	14.91		
NT2RM4001682	12.66	12.66	31.6	33.03	26.04	37.07		
NT2RM4001710	6.7	6.7	38.5	40.58	58.41	40.31		
NT2RM4001712	4.06	4.06	7.61	10.19	10.7	9.98	*	+
NT2RM4001714	10.88	10.88	19.37	18.67	19.3	17.65		
NT2RM4001715	10.77	10.77	11.6	13.55	16.86	12.99		
NT2RM4001727	3.41	3.41	5.92	4.83	5.89	7.6		
NT2RM4001731	2.6	2.6	10.72	13.46	11.23	11.73		
NT2RM4001735	12.84	12.84	21.53	22.01	20.88	34.93		
NT2RM4001739	2.46	2.46	7.3	8.13	5.17	7.14		
NT2RM4001741	14.41	14.41	29.88	26.98	27.21	32.35		
NT2RM4001746	3.65	3.65	6.76	6.89	6.5	5.33		
NT2RM4001754	3.16	3.16	4.17	3.39	3.62	3.84		
NT2RM4001757	5.02	5.02	5.78	4.7	6.31	7.97		
NT2RM4001758	1	1	0.76	1.98	0.65	1.46		
NT2RM4001768	4.83	4.83	10.19	8.48	6.91	7.83		
NT2RM4001775	3.23	3.23	2.76	1.9	1.85	1.71	**	-
NT2RM4001776	2.56	2.56	4.77	2.47	2.68	2.69		
NT2RM4001783	2.88	2.88	3.22	3.12	3.48	3.68		
NT2RM4001793	4.67	4.67	11.44	12.02	9.6	10.75		
NT2RM4001810	3.31	3.31	4.46	3.33	3.63	3.11		
NT2RM4001813	3.9	3.9	4.15	4.71	4.19	5.36		
NT2RM4001818	4.06	4.06	11.34	10.43	8.67	10.53		
NT2RM4001819	2.35	2.35	5.6	2.37	3.02	4.58		
NT2RM4001823	1.76	1.76	4.48	2.47	4.04	4.27		
NT2RM4001828	5.01	5.01	11.49	5.67	7.54	7.51		
NT2RM4001835	9.75	9.75	18.65	21.12	16.5	26.55		
NT2RM4001836	3.27	3.27	8.32	3.65	3.54	5.58		
NT2RM4001841	7.94	7.94	15.82	20.15	20.96	23.33	*	+
NT2RM4001842	2.1	2.1	4.44	3.5	3.35	4.85		
NT2RM4001843	5.65	5.65	14.54	13.34	12.25	14.94		
NT2RM4001856	4.42	4.42	7.16	7.65	4.71	16.83		
NT2RM4001858	5.91	5.91	15.86	16.09	17.03	16.93		
NT2RM4001861	2.91	2.91	9.57	6.31	8.66	9.28		
NT2RM4001863	8.06	8.06	9.5	15.16	15.68	11.77	*	+
NT2RM4001865	5.04	5.04	11.25	7.44	10.24	9.03		
NT2RM4001869	5.1	5.1	5.96	5.22	8.45	21.88		
NT2RM4001873	9.62	9.62	18.43	13.33	15.49	19.21		

(1100)

特開2002-191363

2197

2198

NT2RM4001876	2.24	2.24	6.94	3.65	4.39	7.25		
NT2RM4001880	3.6	3.6	8.57	5.13	5.41	7.67		
NT2RM4001885	5.71	5.71	11.11	7.11	6.56	11.98		
NT2RM4001889	10.25	10.25	18.24	16.31	15.85	21.33		
NT2RM4001894	2.61	2.61	6.07	3.58	3.65	3.49		
NT2RM4001897	7.87	7.87	20.24	18.41	20.4	23.46		
NT2RM4001899	3.36	3.36	7.43	4.92	8.19	8.54		
NT2RM4001905	3	3	4.84	3.3	4.53	7.1		
NT2RM4001922	2.55	2.55	6.05	3.97	4.84	5.11		
NT2RM4001930	2.64	2.64	8.9	2.88	6.53	6.38		
NT2RM4001938	2.65	2.65	4.91	5.09	5.65	6.43		
NT2RM4001940	2.73	2.73	6.17	5.91	4.46	5.48		
NT2RM4001942	37.36	37.36	32.02	53.86	59.28	82.77	*	+
NT2RM4001953	4.65	4.65	9.68	5.04	6.79	4.91		
NT2RM4001965	4.96	4.96	8.82	10.18	8.54	8.39		
NT2RM4001966	3	3	5.14	6.3	7.45	8.19	*	+
NT2RM4001969	2.22	2.22	7.29	5.01	2.95	4.5		
NT2RM4001974	1.19	1.19	4.61	1.89	2.96	4.83		
NT2RM4001979	2.09	2.09	6.37	3.39	4.65	7.36		
NT2RM4001980	4.3	4.3	7.59	7.58	8.02	9.33		
NT2RM4001984	2.31	2.31	5.36	2.68	3.49	4.57		
NT2RM4001987	3.36	3.36	9.66	2.92	4.6	5.01		
NT2RM4002013	6.62	6.62	15.13	13.47	17.16	19.8		
NT2RM4002018	2.31	2.31	5.15	4.09	5.53	7.1		
NT2RM4002033	3.19	3.19	8.16	4.91	3.27	5.93		
NT2RM4002034	1.89	1.89	6.19	4.82	4.38	4.03		
NT2RM4002044	7.71	7.71	17.9	18.75	12.3	18.5		
NT2RM4002047	3.88	3.88	5.19	2.68	5.38	9.2		
NT2RM4002054	4.54	4.54	6.97	2.56	4.3	3.89		
NT2RM4002055	13.72	13.72	74.75	60.51	91.27	61.53		
NT2RM4002059	23.73	23.73	31.85	48.05	63.09	52.61	**	+
NT2RM4002061	3.72	3.72	5.32	3.59	3.69	4.81		
NT2RM4002062	1.9	1.9	5.41	3.66	2.84	4.26		
NT2RM4002063	2.21	2.21	8.1	7.64	7.35	3.79		
NT2RM4002066	2.07	2.07	5.29	4.42	6.32	4.07		
NT2RM4002067	2.51	2.51	4.27	3.07	5.19	4.41		
NT2RM4002073	3.73	3.73	7.24	5.51	7.69	5.16		
NT2RM4002074	5.19	5.19	7.35	5.67	7.47	4.49		
NT2RM4002075	5.13	5.13	5.9	3.16	3.18	2.91	**	-
NT2RM4002076	3.13	3.13	3.05	1.94	2.52	1.71	*	-
NT2RM4002078	10.3	10.3	28.06	23.95	20.81	26.64		
NT2RM4002081	10.47	10.47	30.87	19.18	17.8	18.22		
NT2RM4002082	1.25	1.25	3.02	3.85	2.58	1.23		
NT2RM4002093	2.82	2.82	3.9	4.79	4.66	4.79	*	+
NT2RM4002109	4.42	4.42	11.51	13.95	15.12	15.21	*	+
NT2RM4002115	2.86	2.86	4.51	4.81	4.8	2.52		
NT2RM4002118	4.48	4.48	6.14	4.3	4.86	4.27		
NT2RM4002128	3.78	3.78	4.57	2.84	3.31	3.13	*	-
NT2RM4002137	3.96	3.96	8.14	10.27	7.51	8.92		
NT2RM4002139	3.78	3.78	8.98	7.03	7.84	7.87		

(1101)

特開 2002-191363

2199

2200

NT2RM4002140	4.04	4.04	9.45	8.87	7.81	10.17		
NT2RM4002145	5.99	5.99	17.51	25.81	31.07	24.47	*	+
NT2RM4002146	4.51	4.51	8.23	8.56	9	8.92		
NT2RM4002161	2.33	2.33	4.97	1.38	3.15	5.3		
NT2RM4002174	4.86	4.86	8.02	3.12	4.53	6.15		
NT2RM4002178	7.3	7.3	24.43	28.61	33.13	29.27	*	+
NT2RM4002180	3.47	3.47	11.93	9.27	10.02	11.28		
NT2RM4002185	5.94	5.94	35.51	31.59	32.34	31.69		
NT2RM4002189	1.6	1.6	3.24	3.68	5.59	4.91	*	+
NT2RM4002194	9.3	9.3	25.94	37.2	29.64	38.23	*	+
NT2RM4002198	6.09	6.09	7.61	9.37	8.4	10.04	*	+
NT2RM4002205	4.01	4.01	9.05	6.76	7.86	8.76		
NT2RM4002213	5.36	5.36	8.79	8.05	11.99	14.41		
NT2RM4002216	7.35	7.35	12.58	16.58	23.93	18.16	*	+
NT2RM4002226	3.84	3.84	9.71	20.85	16.65	16.5	**	+
NT2RM4002237	4.19	4.19	10.13	10.37	7.64	13.22		
NT2RM4002240	1.96	1.96	3.64	3.73	3.71	7.59		
NT2RM4002251	2.11	2.11	6.2	7.87	5.48	5.17		
NT2RM4002256	4.38	4.38	10.68	10.7	9.46	9.64		
NT2RM4002262	2.85	2.85	6.25	3.34	4.43	9.66		
NT2RM4002266	3.93	3.93	4.76	2.76	3.55	4.47		
NT2RM4002276	11.23	11.23	15.55	16.5	28.25	20.64		
NT2RM4002278	1.89	1.89	4.59	4.33	3.99	5.11		
NT2RM4002281	17.71	17.71	59.08	62.68	51.19	59.89		
NT2RM4002287	2.08	2.08	3.84	2.46	4.21	3.32		
NT2RM4002294	3.19	3.19	6.99	6.28	6.09	8.69		
NT2RM4002298	18.59	18.59	60.14	86.09	89.9	88.75	*	+
NT2RM4002301	3.2	3.2	6.85	4.63	5.94	4.02		
NT2RM4002306	4.71	4.71	8.24	4.99	5.31	4.2		
NT2RM4002323	3.9	3.9	4.06	4.11	4.39	3.11		
NT2RM4002334	11.54	11.54	20.76	17.92	20.72	16.95		
NT2RM4002339	1.78	1.78	3.52	1.33	1.3	1.38		
NT2RM4002344	2.36	2.36	5.74	2.87	3.57	7.92		
NT2RM4002345	3.56	3.56	10.59	5.06	4.63	7.5		
NT2RM4002352	2.04	2.04	7.67	3.99	5.14	3.74		
NT2RM4002362	20.38	20.38	24.92	11.23	14.32	15.17	*	-
NT2RM4002373	2.1	2.1	3.96	3.21	2.55	3.63		
NT2RM4002374	2.28	2.28	4.39	2.29	3.58	4.3		
NT2RM4002376	4.02	4.02	6.03	3.31	2.97	5.52		
NT2RM4002383	2.8	2.8	8.49	4.76	5.79	4.28		
NT2RM4002390	3.03	3.03	6.01	4.06	5.27	7.37		
NT2RM4002398	5.16	5.16	43.18	33.97	50.73	30.41		
NT2RM4002409	2.11	2.11	5.93	3.37	4.29	1.9		
NT2RM4002414	4.73	4.73	6.21	7.37	9.12	14.53		
NT2RM4002438	2.07	2.07	5.28	3.03	4.38	7.18		
NT2RM4002440	2.99	2.99	6.92	5.78	5.32	9.49		
NT2RM4002446	2.23	2.23	6.08	2.95	4.45	5.7		
NT2RM4002450	3.36	3.36	10.01	6.15	7.75	7.24		
NT2RM4002452	2.13	2.13	6.3	3.67	5.15	7.23		
NT2RM4002457	2.68	2.68	4.44	2.66	3.26	4.52		

(1102)

特開 2002-191363

2201

2202

NT2RM4002458	3.06	3.06	5.77	3.32	5.34	4.04		
NT2RM4002460	2.43	2.43	3.68	1.57	2.45	1.43		
NT2RM4002464	5.4	5.4	12.62	14.39	13.72	14.3		
NT2RM4002479	4.66	4.66	6.69	4.91	7.98	11.54		
NT2RM4002482	4.26	4.26	16.18	10.19	11.5	12.2		
NT2RM4002489	6.74	6.74	16.91	8.79	5.81	11.68		
NT2RM4002493	1.35	1.35	3.22	1.96	3.51	2.73		
NT2RM4002499	34.96	34.96	72.9	59.42	52.07	54.3		
NT2RM4002504	5.15	5.15	10.68	10.57	13.51	9.8		
NT2RM4002506	4.77	4.77	9.4	4.93	7.59	8.53		
NT2RM4002510	2.03	2.03	3.27	1.66	2.97	2.48		
NT2RM4002527	1.57	1.57	3.14	1.83	2.31	4.47		
NT2RM4002532	2.45	2.45	7.75	5.88	3.37	5.19		
NT2RM4002534	1.79	1.79	4.8	2.1	3.45	2.94		
NT2RM4002535	2.5	2.5	6.51	5.89	8.1	6.37		
NT2RM4002554	3.29	3.29	5.31	3.31	5.47	3.12		
NT2RM4002558	6.91	6.91	32.57	32.58	41.54	25.61		
NT2RM4002565	5.38	5.38	13.6	8.22	9.85	10.53		
NT2RM4002567	3.34	3.34	5.43	4.21	4.49	7.22		
NT2RM4002571	4.48	4.48	15.61	11.95	16.47	15.24		
NT2RM4002572	5.57	5.57	17.2	13.7	9.59	13.48		
NT2RM4002577	7.76	7.76	15.25	6.59	5.87	5.65		
NT2RM4002583	1.08	1.08	3.58	2.28	4.83	2.44		
NT2RM4002584	1.64	1.64	5.67	3.24	5.74	5.56		
NT2RM4002593	3.29	3.29	5.17	2.75	3.91	4.61		
NT2RM4002594	11.26	11.26	46.5	38.21	55.32	43.58		
NT2RM4002604	4.83	4.83	4.64	1.77	2.03	2.89	**	-
NT2RM4002614	3.48	3.48	3.48	2.52	3.66	2.81		
NT2RM4002616	1.07	1.07	2.73	2.88	2.71	2.38		
NT2RM4002623	1.39	1.39	4.89	3.92	3.72	5.06		
NT2RM4002634	1.41	1.41	4.38	3.42	4.91	2.96		
NT2RM4002636	2.22	2.22	3.93	3.92	4.18	4.12		
NT2RP1000002	8.82	8.82	52.94	75.1	92.89	81.45	*	+
NT2RP1000006	4.68	4.68	6.28	4.25	4.48	2.56		
NT2RP1000015	4.86	4.86	5.27	2.74	1.99	2.28	**	-
NT2RP1000018	5.45	5.45	5	5.55	4.83	4.96		
NT2RP1000034	18.22	18.22	49.95	38.04	30.76	50.07		
NT2RP1000035	1.93	1.93	3.2	5.26	3.23	3.96		
NT2RP1000040	1.77	1.77	3.33	2.93	3.28	4.28		
NT2RP1000042	1.3	1.3	3.44	1.99	3.22	2.38		
NT2RP1000048	3.6	3.6	10.24	7.25	9.9	9		
NT2RP1000050	2.21	2.21	4.71	2.89	4	3.57		
NT2RP1000056	4.03	4.03	3.74	1.09	0.61	1.96	**	-
NT2RP1000058	3.49	3.49	2.03	1.84	2.07	2.48		
NT2RP1000063	1.77	1.77	3.65	4.09	4	3.83		
NT2RP1000068	1.89	1.89	3.99	3.12	3.33	2.43		
NT2RP1000072	22.9	22.9	74.07	82.91	66.26	95.85		
NT2RP1000073	2.18	2.18	2.45	2.68	3.69	3.86	*	+
NT2RP1000078	2.72	2.72	3.17	2.93	2.3	3.13		
NT2RP1000079	4.13	4.13	5.32	3.6	4.48	2.5		

(1103)

特開2002-191363

2203

2204

NT2RP1000080	4.99	4.99	8.13	9.46	12.46	9.46	*	+
NT2RP1000086	4.15	4.15	3.63	1.31	2.1	3.75		
NT2RP1000087	1.3	1.3	4.36	3.51	3.21	3.45		
NT2RP1000089	4.5	4.5	9.98	12.69	11.3	14.93	*	+
NT2RP1000090	45.76	45.76	96.6	94.37	53.44	93.42		
NT2RP1000100	2.17	2.17	4.05	5.23	4.13	3.66		
NT2RP1000101	3.44	3.44	5.22	4.41	2.88	4.81		
NT2RP1000111	3.24	3.24	5.56	4.51	3.9	3.69		
NT2RP1000112	3.29	3.29	4.08	1.85	3.33	3		
NT2RP1000124	5.57	5.57	4.96	3.11	5.73	5.5		
NT2RP1000125	7.28	7.28	19.39	13.69	10.68	16.86		
NT2RP1000129	1.81	1.81	4.35	5.14	3.91	4.27		
NT2RP1000130	2.31	2.31	4.11	5.31	5.62	16.86		
NT2RP1000154	7.5	7.5	15.63	17.16	12.72	16.37		
NT2RP1000163	2.42	2.42	3.51	2.72	2.99	3.59		
NT2RP1000170	3.42	3.42	4.2	4.96	5.17	5.85	*	+
NT2RP1000174	3.5	3.5	3.42	1.3	2.38	2.12	**	-
NT2RP1000181	6.14	6.14	7.22	10.97	14.98	9.38	*	+
NT2RP1000191	1.08	1.08	5.61	4.94	3.59	5.71		
NT2RP1000202	1.06	1.06	1.66	2.02	1.2	2.24		
NT2RP1000239	1.53	1.53	4.1	2.15	0.94	2.07		
NT2RP1000243	2.37	2.37	2.04	1.31	1.14	1.64	**	-
NT2RP1000255	1.94	1.94	3.02	2.11	2.26	1.78		
NT2RP1000259	5.27	5.27	9.55	5.53	6.33	4.29		
NT2RP1000261	2.76	2.76	4.4	2.07	1.64	2.64		
NT2RP1000269	5.16	5.16	5.01	7.7	10.51	7.39	*	+
NT2RP1000271	7.79	7.79	15.88	13.16	15.11	18.48		
NT2RP1000272	7.71	7.71	13.07	10.72	11.74	11.3		
NT2RP1000279	2.19	2.19	5.24	2.23	3.91	2.62		
NT2RP1000290	6.61	6.61	9.02	12.65	13.52	9.92	*	+
NT2RP1000293	6.86	6.86	10.91	9.75	8.45	10.92		
NT2RP1000300	12.42	12.42	11.93	9.96	11.37	10.2	*	-
NT2RP1000324	5.16	5.16	6	4.69	5.92	6.97		
NT2RP1000325	54.42	54.42	101.4	70.46	57.52	78.6		
NT2RP1000326	4.01	4.01	7.67	3.82	4.56	7.85		
NT2RP1000331	12.16	12.16	24.08	12.19	10.5	20.1		
NT2RP1000333	4.18	4.18	7.52	6.66	6.53	6.98		
NT2RP1000336	1.45	1.45	4.45	1.35	3.76	1.78		
NT2RP1000347	3.05	3.05	8.75	7.26	8.31	6.38		
NT2RP1000348	2.11	2.11	4.14	2.76	3	2.75		
NT2RP1000349	2.12	2.12	3.92	3	4.39	4.11		
NT2RP1000353	40.87	40.87	83.5	51.49	47.8	66.02		
NT2RP1000356	39.53	39.53	93.37	50.3	56.48	74.42		
NT2RP1000357	3.89	3.89	9.63	8.43	8.7	8.72		
NT2RP1000358	2.85	2.85	6.11	4.23	3.04	5.09		
NT2RP1000360	11.04	11.04	19.39	12.08	18.42	19.44		
NT2RP1000363	13.09	13.09	15.39	13.13	13.38	10.01		
NT2RP1000376	1.81	1.81	3.8	2.09	1.9	2.24		
NT2RP1000386	118	118	191.31	146.98	187.97	155.47		
NT2RP1000407	0.72	0.72	3.16	0.58	0.89	1.2		

(1104)

特開 2002-191363

2205

2206

NT2RP1000409	2.05	2.05	5.39	2.84	6.59	3.83		
NT2RP1000413	4.78	4.78	8.03	5.86	8.89	10.19		
NT2RP1000416	1.5	1.5	2.01	0.93	3.17	0.7		
NT2RP1000418	2.27	2.27	6.69	5.08	6.67	4.85		
NT2RP1000420	1.77	1.77	5.19	7.32	7.64	3.7		
NT2RP1000434	1.48	1.48	4.39	1.27	3.12	1		
NT2RP1000439	5.02	5.02	9.31	20.62	28.73	24.75	**	+
NT2RP1000443	1.8	1.8	3.46	2.24	1.61	1.63		
NT2RP1000447	2.21	2.21	5.57	2.49	2.87	3.1		
NT2RP1000448	1.39	1.39	3.58	3.09	4.4	1.41		
NT2RP1000451	4.2	4.2	6.37	5.72	7.27	7.04		
NT2RP1000458	15.1	15.1	10.53	19.73	8.72	23.03		
NT2RP1000460	7.55	7.55	13.82	8.76	11.49	8.62		
NT2RP1000465	4.58	4.58	20.97	20.41	19.98	22.46		
NT2RP1000468	3.25	3.25	4.64	3.82	4.1	4.45		
NT2RP1000470	2.38	2.38	5.67	3.99	2.35	3.8		
NT2RP1000477	1.11	1.11	3.81	1.1	0.84	0.83		
NT2RP1000478	4.53	4.53	12.55	19.87	18.75	20.39	*	+
NT2RP1000481	1.23	1.23	3.89	2.48	4.09	1.2		
NT2RP1000493	2.44	2.44	3.8	1.74	3.83	0.87		
NT2RP1000513	13.07	13.07	16.37	17.06	17.57	18.97	*	+
NT2RP1000522	6.13	6.13	12.69	13.13	13.08	10.32		
NT2RP1000533	3.72	3.72	6.17	2.92	4.49	2.17		
NT2RP1000544	1.53	1.53	2.45	1.38	1.24	1.44		
NT2RP1000547	0.88	0.88	2.45	2	1.63	1.23		
NT2RP1000551	1.7	1.7	2.62	2.13	3.2	1.1		
NT2RP1000567	1.66	1.66	4.29	2.54	4.29	1.77		
NT2RP1000574	1.99	1.99	4.28	1.5	3.43	1.38		
NT2RP1000577	3.14	3.14	6.01	3.16	5.31	2.05		
NT2RP1000579	4.64	4.64	6.24	3.27	3.97	2.04		
NT2RP1000581	5.22	5.22	3.58	2.07	1.61	0.93	**	-
NT2RP1000593	1.74	1.74	4.39	2.48	3.28	2.3		
NT2RP1000604	3.85	3.85	7.75	17.25	13.78	16.39	**	+
NT2RP1000609	1.15	1.15	2.21	2.84	2.61	1.55		
NT2RP1000613	1.12	1.12	2.56	1.82	4.29	0.82		
NT2RP1000622	5.94	5.94	15.9	14.91	19.42	15.46		
NT2RP1000627	9.18	9.18	18.96	23.88	21.9	14.86		
NT2RP1000629	4.18	4.18	5.9	5.92	5.32	3.17		
NT2RP1000630	6.54	6.54	7.84	7.21	7.67	7.92		
NT2RP1000639	0.64	0.64	0.31	1.53	2.04	0.28		
NT2RP1000640	130.14	130.14	307.77	227.5	176.05	232.29		
NT2RP1000646	4.14	4.14	9.59	10.19	11.87	12.15	*	+
NT2RP1000659	2.65	2.65	7	8.91	7.99	6.04		
NT2RP1000674	13.48	13.48	28.08	43.62	45.82	56.95	**	+
NT2RP1000677	3.9	3.9	10.76	11.84	10.19	9.87		
NT2RP1000679	2.38	2.38	3.76	2.3	2.35	1.05		
NT2RP1000688	4.72	4.72	3.34	2.76	2.73	1.83	*	-
NT2RP1000689	1.44	1.44	1.86	2.03	1.22	1.13		
NT2RP1000695	1.11	1.11	2.5	2.09	2.44	1.52		
NT2RP1000701	0.89	0.89	1.08	2.62	2.74	1.71	*	+

(1105)

特開2002-191363

2207

2208

NT2RP1000702	1.12	1.12	2.28	3.74	4.07	3.15	*	+
NT2RP1000713	2.29	2.29	2.79	2.8	3.56	2.38		
NT2RP1000721	4.14	4.14	4.49	4.48	3.92	3.78		
NT2RP1000730	3.5	3.5	4.83	2.61	4.5	2.41		
NT2RP1000733	6.08	6.08	6.56	4.91	8.12	5.65		
NT2RP1000738	3.18	3.18	8.04	5.16	5.71	7.11		
NT2RP1000739	1.11	1.11	2.65	4.02	3.09	2.86		
NT2RP1000740	1.41	1.41	3.13	3.63	3.57	3.77	*	+
NT2RP1000746	1.15	1.15	3.58	2.28	3.74	1.37		
NT2RP1000750	4	4	8.31	10.25	10.72	9.39	*	+
NT2RP1000751	33.15	33.15	59.65	67.84	64.22	66.55		
NT2RP1000767	3.8	3.8	3.64	1.7	2.62	0.62	*	-
NT2RP1000769	9.31	9.31	13.98	7.42	8.59	7.19		
NT2RP1000780	0.86	0.86	1.01	1.87	1.13	0.89		
NT2RP1000782	4.25	4.25	12.21	10.24	7.96	9.13		
NT2RP1000796	3.17	3.17	2.69	4.23	2.99	2.86		
NT2RP1000797	12.31	12.31	22.78	19.44	18.64	21.78		
NT2RP1000800	1.13	1.13	3.74	2.46	2.66	1.46		
NT2RP1000825	2.38	2.38	2.91	1.04	1.88	0.87	*	-
NT2RP1000833	2.5	2.5	2.85	0.92	2.39	1.26		
NT2RP1000834	35.44	35.44	66.57	73.98	90.28	71.45		
NT2RP1000836	1.83	1.83	3.43	1.01	2.39	1.04		
NT2RP1000837	3.36	3.36	6.66	3.22	4.71	3.67		
NT2RP1000846	1.29	1.29	5.48	1.67	2.84	1.4		
NT2RP1000847	1.99	1.99	5.49	2.15	5.12	1.64		
NT2RP1000851	4.67	4.67	9.32	6.18	7.94	6.72		
NT2RP1000856	14.31	14.31	17.46	20.38	23.22	19.37	*	+
NT2RP1000860	2.09	2.09	4.54	4.02	2.74	4.04		
NT2RP1000902	5.31	5.31	11.6	6.94	9.91	7.34		
NT2RP1000903	2.45	2.45	6.26	4.04	3.42	4.24		
NT2RP1000905	1.76	1.76	4.87	5.36	5.66	10		
NT2RP1000915	5.51	5.51	10.01	6.72	8.59	9.91		
NT2RP1000916	2.31	2.31	5.51	1.78	3.82	2.09		
NT2RP1000921	9.38	9.38	8.73	8.23	9.13	7.92		
NT2RP1000943	5.14	5.14	10.76	8.51	8.55	7.2		
NT2RP1000944	1.59	1.59	2.21	1.78	1.74	1.15		
NT2RP1000947	8.5	8.5	14.91	16.51	15.04	14.22		
NT2RP1000954	2.11	2.11	4.96	2.74	5.55	3.04		
NT2RP1000958	6.48	6.48	14.73	4.54	10.17	10.21		
NT2RP1000959	124.81	124.81	209.45	128.43	72.65	206.1		
NT2RP1000966	9.96	9.96	12.96	14.28	15.36	21.39		
NT2RP1000974	2.46	2.46	5.38	3.98	6.08	3.71		
NT2RP1000980	3.07	3.07	5.5	4.04	4.53	4.02		
NT2RP1000981	4.3	4.3	8.09	5.68	7.26	5.27		
NT2RP1000988	6.45	6.45	10.46	9.62	6.44	7.87		
NT2RP1001002	2.8	2.8	7.36	3.94	4.57	4.3		
NT2RP1001004	4.72	4.72	8.25	3.65	4.9	5.37		
NT2RP1001007	1.42	1.42	3.42	1.69	3.84	2.03		
NT2RP1001011	1.94	1.94	5.93	3.82	5.46	4.83		
NT2RP1001013	4.45	4.45	9.41	5.92	8.62	5.04		

(1106)

特開2002-191363

2209

2210

NT2RP1001014	2.21	2.21	5.89	3.76	6.64	3.49		
NT2RP1001020	1.87	1.87	4.11	2.08	3.75	2.36		
NT2RP1001023	62.79	62.79	145.09	101.48	105.86	143.96		
NT2RP1001027	18.11	18.11	82.66	51.63	59.05	68.9		
NT2RP1001031	1.83	1.83	3.31	2.05	3.53	2.25		
NT2RP1001033	2.43	2.43	6.09	5.68	5.98	4.27		
NT2RP1001042	2.94	2.94	6.47	2.99	3.85	2.04		
NT2RP1001045	15.95	15.95	23.24	40.66	47.04	44.53	**	+
NT2RP1001073	6.64	6.64	10.57	8.32	10.46	7.33		
NT2RP1001079	2.91	2.91	6.37	2.16	2.58	1.48		
NT2RP1001080	2.16	2.16	4.89	6.88	4.2	4.56		
NT2RP1001113	1.07	1.07	3.64	3.55	3.94	3.26		
NT2RP1001159	21.42	21.42	43.84	22.89	23.31	34.25		
NT2RP1001173	1.7	1.7	3.07	1.38	4.28	1.52		
NT2RP1001176	7.4	7.4	10.13	13	9.31	13.95		
NT2RP1001177	5.31	5.31	5.75	3.01	5.5	2.02		
NT2RP1001185	6.42	6.42	9.37	3.79	4.63	2.73	*	-
NT2RP1001199	3.9	3.9	7.67	6.93	5.22	3.28		
NT2RP1001205	7.78	7.78	19.46	16.66	12.64	23.28		
NT2RP1001215	1.82	1.82	5.02	3.79	4.12	3.15		
NT2RP1001225	4.54	4.54	7.96	7.56	8.77	6.31		
NT2RP1001245	7.27	7.27	10.86	19.68	21.03	22.13	**	+
NT2RP1001247	2.04	2.04	4.01	1.77	2.89	1.67		
NT2RP1001248	2.81	2.81	6.79	3.94	4.63	2.4		
NT2RP1001253	5.02	5.02	6.39	4.48	4.38	3.32		
NT2RP1001286	6.18	6.18	7.69	3.79	3.88	4.12	**	-
NT2RP1001294	2.4	2.4	4.47	3.6	2.73	4.18		
NT2RP1001302	2.46	2.46	4.51	4.89	2.9	5.39		
NT2RP1001310	15.54	15.54	34.01	21.13	20.75	27.15		
NT2RP1001311	1.9	1.9	3.22	2.66	3.16	2.38		
NT2RP1001313	2.6	2.6	7.72	5.45	7.85	5.78		
NT2RP1001324	2.47	2.47	5.3	3.34	4.17	2.35		
NT2RP1001349	3.3	3.3	6.29	3.63	3.92	2.14		
NT2RP1001361	19.41	19.41	18.28	23.28	28.33	24.16	*	+
NT2RP1001379	3.82	3.82	9.52	4.97	7.97	7.06		
NT2RP1001385	2.06	2.06	4.51	4.09	3.89	4.4		
NT2RP1001395	4.96	4.96	7.86	6.01	6.32	8.13		
NT2RP1001410	8.75	8.75	20.39	15.74	15.66	9.94		
NT2RP1001424	2.39	2.39	3.34	3	3	1.73		
NT2RP1001432	4.33	4.33	3.86	2.19	1.76	2.05	**	-
NT2RP1001449	6.23	6.23	7.5	6.29	8.21	4.63		
NT2RP1001457	4.09	4.09	4.21	2.11	2.26	2.63	**	-
NT2RP1001459	21.54	21.54	132.97	90.96	107.97	81.08		
NT2RP1001466	5.73	5.73	14.97	11.31	9.39	10.99		
NT2RP1001475	2.45	2.45	6.31	5.98	6.67	3.9		
NT2RP1001482	3.93	3.93	9.18	15.88	13.03	8.2		
NT2RP1001494	1.61	1.61	4.6	4.34	4.18	2.25		
NT2RP1001500	3.39	3.39	8.13	8.09	8.65	7.42		
NT2RP1001517	5.11	5.11	7.37	4.41	5.38	2.36		
NT2RP1001540	4.74	4.74	5.03	4.6	4.86	3.11		

(1107)

特開 2002-191363

2211

2212

NT2RP1001543	1.02	1.02	1.83	1.49	1.12	0.98		
NT2RP1001546	22.51	22.51	51.51	34.99	22.76	33.42		
NT2RP1001550	9.33	9.33	21.4	14.35	12.21	13.42		
NT2RP1001553	2.07	2.07	6.07	5.69	6.04	4.45		
NT2RP1001555	36.28	36.28	58.55	41.1	53.63	54.35		
NT2RP1001563	2.28	2.28	3.44	2.07	2.24	1.31		
NT2RP1001569	9.43	9.43	16	17.31	18.21	13.04		
NT2RP1001584	15.6	15.6	19.66	28.1	32.53	25.83	**	+
NT2RP1001599	1.18	1.18	1.95	1.27	1.19	1.24		
NT2RP1001616	5	5	11.95	9.49	6.7	8.95		
NT2RP1001654	11.78	11.78	18.07	16.27	16.54	18.38		
NT2RP1001665	2.77	2.77	4.72	2.73	2.05	2.05		
NT2RP1001679	76.31	76.31	195.7	199.2	240.87	222.46		
NT2RP1001681	10.11	10.11	15.1	20.39	21.03	23.99	**	+
NT2RP1001694	3.58	3.58	3.82	2.45	2.38	1.97	**	-
NT2RP2000001	5.23	5.23	5.53	4.17	4.54	3.74	**	-
NT2RP2000006	3.49	3.49	7.32	4.12	3.88	3.4		
NT2RP2000007	3.18	3.18	6.56	4.68	5.66	4.92		
NT2RP2000008	2.77	2.77	6.72	3.66	5.3	4.9		
NT2RP2000010	2.89	2.89	5.59	2.99	5.06	2.5		
NT2RP2000011	7.08	7.08	17.96	14.55	14.74	15.15		
NT2RP2000027	2.28	2.28	7.42	4.52	4.89	3.61		
NT2RP2000028	22.93	22.93	62.54	46.48	51.47	53.47		
NT2RP2000032	2.5	2.5	5.85	3.11	3.71	6.42		
NT2RP2000040	11.57	11.57	23.92	14.38	14.5	23.1		
NT2RP2000042	5.28	5.28	10.32	6.89	7.21	12.64		
NT2RP2000045	5.7	5.7	9.42	5.27	6.45	6.3		
NT2RP2000051	3.16	3.16	6.29	9.23	9.96	9.53	**	+
NT2RP2000054	2.55	2.55	6.42	3.81	5.42	2.53		
NT2RP2000056	3.68	3.68	6.23	5.67	6.89	5.8		
NT2RP2000057	60.79	60.79	174.83	212.63	239.81	221.98	*	+
NT2RP2000067	3.1	3.1	3.86	2.98	4.36	5.72		
NT2RP2000070	2.91	2.91	6.27	5.7	5.95	8.21		
NT2RP2000076	1.66	1.66	4.45	2.98	3.58	3.23		
NT2RP2000077	1.67	1.67	4.73	2.43	4.94	3.14		
NT2RP2000079	3.76	3.76	9.24	5.15	4.81	5.47		
NT2RP2000088	2.9	2.9	5.22	2.18	3.07	2.21		
NT2RP2000091	5.84	5.84	6.54	6.62	8.28	6.72		
NT2RP2000092	4.37	4.37	6.7	6.06	7.67	5.65		
NT2RP2000097	2.74	2.74	3.39	3.4	4.13	4.13	*	+
NT2RP2000098	3.44	3.44	6.83	6.69	9.01	6.27		
NT2RP2000108	1.93	1.93	7.24	4.8	6.31	6.68		
NT2RP2000114	1.95	1.95	3.65	2.58	4.41	2.9		
NT2RP2000116	3.17	3.17	7.36	5.35	3.85	9.42		
NT2RP2000119	3.14	3.14	7.16	4.58	7.96	5.6		
NT2RP2000120	3.91	3.91	7.62	5.57	8.5	5.8		
NT2RP2000126	2.86	2.86	4.86	3.88	5.1	3.44		
NT2RP2000133	1.83	1.83	3.66	3.13	4.05	2.01		
NT2RP2000147	6.28	6.28	12.88	11.64	6.51	8.58		
NT2RP2000153	4.61	4.61	9.55	10.57	6.49	12.05		

(1108)

特開 2002-191363

2213

2214

NT2RP2000156	3.27	3.27	8.24	5.59	6.55	4.07		
NT2RP2000157	3.7	3.7	6.33	6.57	5.02	4.15		
NT2RP2000161	4.45	4.45	8.82	7.52	7.5	6.02		
NT2RP2000168	4.22	4.22	12.63	3.94	6.03	3.88		
NT2RP2000173	12.56	12.56	81.37	72.12	92.12	78.68		
NT2RP2000175	1.9	1.9	3.78	2.03	3.33	4.43		
NT2RP2000178	2.06	2.06	6.06	3.23	3.01	3.81		
NT2RP2000183	1.64	1.64	7.82	5.82	6.21	6.03		
NT2RP2000195	3.1	3.1	6.65	6	6.71	3.22		
NT2RP2000204	73.6	73.6	93.43	102.95	40.16	62.34		
NT2RP2000205	4	4	6.56	3.91	5.74	4.5		
NT2RP2000208	3.06	3.06	9.42	4.23	6.77	3.36		
NT2RP2000224	13.3	13.3	31.75	18.34	21.15	20.93		
NT2RP2000230	9.96	9.96	18.99	12.16	16.4	11.95		
NT2RP2000231	4.3	4.3	7.41	4.24	3.43	4.54		
NT2RP2000232	1.08	1.08	2.75	1.53	2.45	0.74		
NT2RP2000233	8.04	8.04	60.44	47.2	64.72	52.31		
NT2RP2000239	3	3	4.7	8.93	8.93	7.01	**	+
NT2RP2000240	2.01	2.01	5.25	2.49	3.41	1.45		
NT2RP2000248	4.29	4.29	6.09	2.82	2.39	0.96	*	-
NT2RP2000256	5.7	5.7	8.25	5.62	6.22	5.44		
NT2RP2000257	3.47	3.47	6.92	4.86	6.52	4.34		
NT2RP2000258	1.53	1.53	3.83	3.88	2.93	3.02		
NT2RP2000261	2.95	2.95	3.94	4.47	3.59	2.91		
NT2RP2000270	3.12	3.12	6.26	6.66	4.06	4.3		
NT2RP2000274	1.78	1.78	3.87	3.48	5.16	2.56		
NT2RP2000277	2.18	2.18	6.13	3.19	4.02	2.98		
NT2RP2000279	2.26	2.26	4.92	2.43	2.52	2.17		
NT2RP2000283	5.75	5.75	27.65	21.53	27.32	24.78		
NT2RP2000288	6.29	6.29	7.46	8.9	10.89	8.64	*	+
NT2RP2000289	1.12	1.12	2.79	3.09	2.77	1.93		
NT2RP2000297	2.57	2.57	5.7	4.8	4.53	6.09		
NT2RP2000298	3.61	3.61	9.64	8.51	7.66	8.24		
NT2RP2000310	1.43	1.43	2.3	2.46	3.41	1.7		
NT2RP2000327	2.12	2.12	3.96	3.13	3.49	1.57		
NT2RP2000328	6.95	6.95	11.56	13.43	16.7	14.68	*	+
NT2RP2000329	10.73	10.73	10.17	17.55	23.92	18.52	**	+
NT2RP2000333	6.35	6.35	6.4	6.83	7.17	4.64		
NT2RP2000337	2.05	2.05	5.16	4.43	5.32	5.31		
NT2RP2000346	2.55	2.55	5.18	7.2	5.63	4.95		
NT2RP2000357	1.57	1.57	6.87	5.48	5.14	5.35		
NT2RP2000358	2.09	2.09	4.52	5.03	4.9	4.01		
NT2RP2000366	3.23	3.23	4.08	4.16	4.25	2.32		
NT2RP2000369	7.22	7.22	9.94	44.13	45.2	44.34	**	+
NT2RP2000376	26.92	26.92	108.62	84.48	134.63	85.95		
NT2RP2000394	6.49	6.49	5.92	5.08	8.52	4.21		
NT2RP2000396	2.71	2.71	6.55	7.52	6.8	5.02		
NT2RP2000412	4.48	4.48	23.45	21.42	24.93	20.49		
NT2RP2000414	8.03	8.03	18.69	23.83	18.98	23.37		
NT2RP2000420	1.12	1.12	4.11	3.54	3.25	1.97		

(1109)

特開2002-191363

2215

2216

NT2RP2000422	6.41	6.41	13.18	17.56	17.88	18.67	*	+
NT2RP2000426	21.59	21.59	80.94	87.94	110.97	74.98		
NT2RP2000428	24.92	24.92	43.91	34.21	35.59	30.95		
NT2RP2000438	5.06	5.06	5.17	5.62	6.94	5.11		
NT2RP2000447	4.14	4.14	9.68	7.3	7.08	7.16		
NT2RP2000448	3.03	3.03	4.63	4.57	3.57	3.17		
NT2RP2000459	2.47	2.47	4.93	2.82	3.15	2.09		
NT2RP2000479	3.3	3.3	7.51	5.33	5.71	5.06		
NT2RP2000498	3.07	3.07	6.25	4.48	5.09	3.9		
NT2RP2000503	2.47	2.47	4.46	2.54	2.82	1.52		
NT2RP2000510	4.01	4.01	6.19	5.08	6.45	3.7		
NT2RP2000514	2.65	2.65	2.51	1.94	2.25	1.63	*	-
NT2RP2000516	4.72	4.72	9.77	4.92	5.29	5.18		
NT2RP2000523	2.21	2.21	3.17	1.92	2.44	2.63		
NT2RP2000533	17.82	17.82	29.05	22.57	27.56	30.78		
NT2RP2000540	1.98	1.98	4.66	3.01	5.41	5.18		
NT2RP2000547	3.1	3.1	5.26	4.38	5.27	3.71		
NT2RP2000557	4.26	4.26	6.96	4.34	6.5	3.32		
NT2RP2000558	3.43	3.43	7.17	6.43	7.11	8.26		
NT2RP2000564	3.04	3.04	7.2	3.49	8.03	4.77		
NT2RP2000565	4.54	4.54	11.07	7.64	9.24	9.98		
NT2RP2000583	14.8	14.8	44.9	49.6	34.93	49.08		
NT2RP2000591	0.81	0.81	3.81	1.53	2.61	1.21		
NT2RP2000599	1.85	1.85	4.1	1.97	3.43	2.36		
NT2RP2000601	1.78	1.78	4.67	1.28	2.48	1.3		
NT2RP2000603	2.58	2.58	4.44	2.54	2.84	2.98		
NT2RP2000610	3.77	3.77	7.23	6.32	7.62	5.53		
NT2RP2000614	75.85	75.85	129.42	130.63	184.38	188.58	*	+
NT2RP2000616	1.81	1.81	4.89	3.9	5.1	3.83		
NT2RP2000617	2.17	2.17	6.73	5.78	6.82	6.26		
NT2RP2000623	3.1	3.1	5.36	3.46	5.1	3.49		
NT2RP2000634	1.56	1.56	3.92	2.29	3.34	2.02		
NT2RP2000636	3.78	3.78	8.64	6.27	7.6	6.62		
NT2RP2000638	4.37	4.37	8.91	4.57	7.41	5.69		
NT2RP2000644	2.22	2.22	5.47	3.41	4.16	3.45		
NT2RP2000649	8.96	8.96	15.76	13.65	17.22	13.07		
NT2RP2000652	3.35	3.35	4.58	3.57	4.36	2.72		
NT2RP2000656	3.73	3.73	6.93	4.83	3.91	4.08		
NT2RP2000658	1.08	1.08	2.64	1.51	3.18	1.43		
NT2RP2000663	4.23	4.23	6.9	5.98	7.21	5.9		
NT2RP2000664	4.24	4.24	10.24	12.72	12.54	16.44	*	+
NT2RP2000668	7.49	7.49	26.84	16.92	20.41	17.17		
NT2RP2000678	1.77	1.77	3.19	1.77	2.09	1.13		
NT2RP2000694	4.89	4.89	8.39	11.06	13	13.36	**	+
NT2RP2000704	1.8	1.8	5.63	2.99	3.13	3.67		
NT2RP2000710	4.51	4.51	9.96	6.72	8.08	7.23		
NT2RP2000712	1.43	1.43	5.35	3.39	3.57	2.95		
NT2RP2000715	3.42	3.42	7.43	6.04	7.56	4.49		
NT2RP2000720	4.92	4.92	11.76	7.24	8	7.11		
NT2RP2000731	3.92	3.92	9.15	3.7	4.5	2.61		

(1110)

特開 2002-191363

2217

2218

NT2RP2000739	3.23	3.23	5.67	2.62	3.32	5.65		
NT2RP2000748	1.59	1.59	4.2	1.42	1.81	1.62		
NT2RP2000749	11.84	11.84	21.88	14.4	8.47	13.91		
NT2RP2000758	1.6	1.6	3.17	2.65	6	1.17		
NT2RP2000764	1.51	1.51	5.74	2.95	5.22	1.95		
NT2RP2000766	9.08	9.08	52.24	46.37	59.37	52.89		
NT2RP2000777	12.28	12.28	18.43	26.91	28.56	24.47	**	+
NT2RP2000786	21.32	21.32	73.91	55.85	67.59	58.16		
NT2RP2000793	5.32	5.32	6.9	4.32	3.57	4.38	*	-
NT2RP2000796	5.32	5.32	7.41	7.38	9.17	6.66		
NT2RP2000809	3.25	3.25	8.3	6.46	4.69	5.45		
NT2RP2000812	6.65	6.65	17.51	16.43	14.35	16.89		
NT2RP2000814	4.16	4.16	4.97	3.75	4.6	3.29		
NT2RP2000816	1.84	1.84	5.64	4.64	5.19	3.58		
NT2RP2000818	3.28	3.28	5.19	3.18	3.66	1.95		
NT2RP2000819	2.76	2.76	5.79	3.03	3.05	1.94		
NT2RP2000841	4.35	4.35	4.51	2.17	2.48	1.65	**	-
NT2RP2000842	7.8	7.8	9.57	13.62	14.25	12.66	**	+
NT2RP2000845	2.52	2.52	8.31	7.51	6.76	6.93		
NT2RP2000863	2.45	2.45	3.48	3.82	3.37	2.47		
NT2RP2000880	5.96	5.96	11.61	9.5	11.13	10.25		
NT2RP2000892	4.3	4.3	6.43	6.54	6.97	5.01		
NT2RP2000894	5.59	5.59	11.88	5.41	5.59	2.16		
NT2RP2000903	5.71	5.71	9.12	10.73	11.92	7.44		
NT2RP2000906	4.56	4.56	5.39	2.63	3.78	2.19	*	-
NT2RP2000910	4.34	4.34	4.26	2.9	2.7	1.68	**	-
NT2RP2000931	10.97	10.97	18.36	20.51	19.28	24.6		
NT2RP2000932	2.86	2.86	5.43	4.8	4.72	4.21		
NT2RP2000938	18.41	18.41	42.99	35.71	30.01	43.52		
NT2RP2000943	7.02	7.02	14.98	18.7	14.88	14.48		
NT2RP2000957	3.19	3.19	4.11	4.26	3.66	2.71		
NT2RP2000958	7	7	6.84	10.43	12.36	7.8		
NT2RP2000959	9.88	9.88	14.99	13.92	17.38	10.69		
NT2RP2000965	5.05	5.05	7.82	15.73	18.97	16.02	**	+
NT2RP2000970	2.31	2.31	6.72	5.14	5	4.62		
NT2RP2000973	0.9	0.9	1.47	2.56	2.64	1.57	*	+
NT2RP2000985	2.69	2.69	6	9.28	6.29	13.98		
NT2RP2000987	1.89	1.89	3.31	4.54	3.17	1.66		
NT2RP2000997	13.83	13.83	23.99	38.12	29.73	40.96	*	+
NT2RP2001024	2.86	2.86	5.34	3.61	3.12	2.36		
NT2RP2001028	4.66	4.66	4.2	2.65	4.09	0.99		
NT2RP2001036	5.14	5.14	8.86	6.16	6.44	4.91		
NT2RP2001039	1.08	1.08	3.18	3.47	1.14	2.24		
NT2RP2001044	1.13	1.13	2.5	2.53	1.89	2.8		
NT2RP2001056	4.97	4.97	28.32	20.16	26.9	18.16		
NT2RP2001065	2.38	2.38	6.24	7.45	7.4	5.69		
NT2RP2001067	2.98	2.98	5.38	4.12	5.29	2.41		
NT2RP2001070	3.3	3.3	7.63	4.72	6.17	3.58		
NT2RP2001081	2.91	2.91	8.19	4.8	6.68	3.9		
NT2RP2001087	3.93	3.93	2.36	2.06	2.92	1.61		

(1111)

特開 2002-191363

2219

2220

NT2RP2001094	0.69	0.69	1.37	1.25	1.15	1.04
NT2RP2001119	2.02	2.02	6.11	5.86	4.44	4.35
NT2RP2001127	1.53	1.53	4.04	2.69	1.85	2.1
NT2RP2001133	2.45	2.45	4.73	4.06	3.61	3.6
NT2RP2001137	2.68	2.68	4.07	2.3	2.82	2.9
NT2RP2001142	3.88	3.88	7.47	3.37	2.83	2.42
NT2RP2001149	2.7	2.7	2.98	2.11	3.39	1.32
NT2RP2001168	6	6	7.81	6.8	7.01	5.75
NT2RP2001173	4.15	4.15	7.88	3.98	3.09	5.44
NT2RP2001174	9.23	9.23	14.98	14.12	15.45	18.01
NT2RP2001184	2.78	2.78	5.46	4.21	7.18	4.64
NT2RP2001196	1.62	1.62	5.93	3.14	3.28	2.94
NT2RP2001200	3.85	3.85	9.36	5.02	4.25	7.88
NT2RP2001218	2.29	2.29	5.69	2.55	3.89	3.6
NT2RP2001223	2.65	2.65	5.03	1.95	3.69	3.31
NT2RP2001226	4.34	4.34	10.19	6.95	5.72	7.35
NT2RP2001227	4.45	4.45	6.12	3.2	3.62	6.01
NT2RP2001232	6.44	6.44	13.95	7.13	9.79	13.66
NT2RP2001233	4.02	4.02	10.57	7.04	7.77	8.01
NT2RP2001245	4.21	4.21	8.03	9.47	11.82	11.16
NT2RP2001246	6.3	6.3	9.84	9	11.28	11.57
NT2RP2001268	6.19	6.19	18.1	17.61	16.26	18.55
NT2RP2001270	4.78	4.78	9.11	5.68	8.7	10.04
NT2RP2001276	4.92	4.92	13.29	12.73	10.92	12.73
NT2RP2001277	3.11	3.11	7.02	4.91	6.22	10.82
NT2RP2001290	2.71	2.71	6.46	4.42	5.61	6.01
NT2RP2001295	5.46	5.46	9.44	5.13	5.98	7.92
NT2RP2001297	118.17	118.17	120.73	139.11	97.16	145.76
NT2RP2001301	9.12	9.12	18.56	15.89	19.62	14.24
NT2RP2001312	2.7	2.7	5.68	5.6	4.59	6.04
NT2RP2001327	4.73	4.73	5.69	6.39	8.53	11.86
NT2RP2001328	8.44	8.44	20.87	16.32	23.25	23.16
NT2RP2001341	4.59	4.59	9.22	3.06	7.65	7.21
NT2RP2001347	3.09	3.09	8.54	5.54	9.55	6.9
NT2RP2001366	10.33	10.33	48.06	54.83	51.5	52.33
NT2RP2001378	2.33	2.33	3.77	3.74	4.64	5.02
NT2RP2001381	2.82	2.82	6.86	5.79	6.62	8.37
NT2RP2001388	3.25	3.25	6.71	4.54	5.11	5.2
NT2RP2001391	443.52	443.52	734.13	742.83	990.71	747.95
NT2RP2001392	2.98	2.98	6.43	4.58	3.16	4.18
NT2RP2001394	3.3	3.3	8.55	8.35	6.09	10.15
NT2RP2001397	5.04	5.04	6.79	7.33	5.68	12.2
NT2RP2001400	3.1	3.1	6.4	3.43	6.25	2.92
NT2RP2001408	3.31	3.31	6.13	4.02	5.97	5.62
NT2RP2001420	5.63	5.63	12.09	8.09	9.97	9.17
NT2RP2001423	4.71	4.71	9.71	6.21	8.29	7.19
NT2RP2001427	2.68	2.68	5.32	3.69	4.61	5.49
NT2RP2001428	2.71	2.71	7.13	5.49	3.78	3.03
NT2RP2001436	4.27	4.27	8.85	5.84	2.85	4.84
NT2RP2001440	2.89	2.89	7.34	10.24	10.15	11.98

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(1112)

特開2002-191363

2221

2222

NT2RP2001445	2.43	2.43	6.75	5.86	5.55	5.89		
NT2RP2001449	4.37	4.37	6.41	5	4.74	5.02		
NT2RP2001450	3.19	3.19	6.75	2.26	5.4	8.59		
NT2RP2001467	4.53	4.53	10.28	5.32	4.72	6.5		
NT2RP2001469	4.74	4.74	6.79	8.22	11.04	7.18		
NT2RP2001480	6.54	6.54	26.68	14.98	12.63	15.42		
NT2RP2001495	5.86	5.86	11.96	8.16	9.04	10.39		
NT2RP2001499	8.25	8.25	16.78	10.05	14.46	9.66		
NT2RP2001506	2.79	2.79	7.24	5.32	8.19	5.33		
NT2RP2001508	10.59	10.59	13.66	18.74	20.49	21.92	**	+
NT2RP2001511	6.41	6.41	9.74	6.08	8.63	6.53		
NT2RP2001514	7.04	7.04	7.02	7.24	6.44	6.38		
NT2RP2001520	2.93	2.93	4.84	2.6	3.19	2.87		
NT2RP2001526	3.88	3.88	8.49	7.01	5.27	3.83		
NT2RP2001529	9.87	9.87	53.78	44.74	55.72	60.88		
NT2RP2001536	1.63	1.63	4.17	3.81	4.52	4.71		
NT2RP2001538	83.44	83.44	178.68	132.75	146.73	155.87		
NT2RP2001547	4.96	4.96	14.87	16.77	19.21	17.77	*	+
NT2RP2001560	6.28	6.28	21.64	25.41	28.19	27.75	*	+
NT2RP2001562	5.56	5.56	5.57	5.94	6.75	4.64		
NT2RP2001566	7.96	7.96	9.24	8.22	7.91	8.56		
NT2RP2001569	4.26	4.26	8.71	6.09	5.65	8.28		
NT2RP2001576	3.95	3.95	11.58	13.42	9	12.82		
NT2RP2001581	47.15	47.15	130.15	121.19	112.28	129.54		
NT2RP2001597	3.73	3.73	7.88	8.57	8.3	13.3		
NT2RP2001601	2.37	2.37	4.81	3.67	4.81	3.34		
NT2RP2001613	2.74	2.74	4.87	2.83	2.72	4.15		
NT2RP2001628	3.42	3.42	3.97	3.14	3.04	7.84		
NT2RP2001634	8.64	8.64	13.94	16.57	23.67	17.67	*	+
NT2RP2001635	2.51	2.51	5.92	5.63	5.53	4.72		
NT2RP2001660	4.27	4.27	16.91	5.9	5.54	10.06		
NT2RP2001662	1.49	1.49	4.07	4.5	4.44	3.47		
NT2RP2001663	2.82	2.82	5.09	10.37	8.21	9.74	**	+
NT2RP2001672	3.28	3.28	3.82	3.88	4.09	4	*	+
NT2RP2001675	4.1	4.1	5.01	5.23	4.73	5.06		
NT2RP2001677	9.58	9.58	18.2	20.9	26.67	19.74	*	+
NT2RP2001678	4.84	4.84	6.73	4.6	4.83	4.5		
NT2RP2001683	1.89	1.89	3.12	4.6	4.72	2.78		
NT2RP2001699	3.15	3.15	6.16	6.5	5.84	4.88		
NT2RP2001707	1.24	1.24	3.19	3.42	4.13	4.8	*	+
NT2RP2001720	1.47	1.47	3.6	3.91	3	2.72		
NT2RP2001721	2.26	2.26	4.57	5.53	3.96	3.66		
NT2RP2001740	12	12	60.21	52.38	79.71	54.73		
NT2RP2001748	6.43	6.43	10.8	8.75	10.25	8.55		
NT2RP2001755	5.51	5.51	4.96	3.71	4.62	2.69	*	-
NT2RP2001762	1.25	1.25	2.01	3.87	2.56	3.52	*	+
NT2RP2001768	1.91	1.91	4.7	6.7	5.55	4.55		
NT2RP2001769	3.06	3.06	5.86	10.42	5.06	11.86		
NT2RP2001784	3.62	3.62	6.23	7.06	6.02	6.91		
NT2RP2001805	2.33	2.33	5.61	6.02	4.93	6.6		

(1113)

特開 2002-191363

2223

2224

NT2RP2001813	2.75	2.75	3.73	1.84	1.98	1.94	*	-
NT2RP2001817	3.16	3.16	4.49	4.03	5.32	3.45		
NT2RP2001818	2.72	2.72	2.45	2.35	3.62	2.66		
NT2RP2001837	5.13	5.13	13.43	10.29	10.16	12.33		
NT2RP2001839	17.02	17.02	83.84	60.14	71.06	82.26		
NT2RP2001861	2	2	6.37	3.16	3.52	3.87		
NT2RP2001869	2.64	2.64	6.54	4.35	5.77	8.84		
NT2RP2001876	12.15	12.15	27.71	24.54	24.93	23.67		
NT2RP2001878	2.32	2.32	3.96	2.95	3.32	4.95		
NT2RP2001881	3.72	3.72	5.4	9.67	12.64	12.16	**	+
NT2RP2001883	2.63	2.63	6.8	4.33	5.42	6.35		
NT2RP2001884	13.59	13.59	23.56	15.33	10.54	23.6		
NT2RP2001885	3.27	3.27	5.49	2.88	4.39	4.82		
NT2RP2001898	10.76	10.76	80.37	69.48	88.43	73.46		
NT2RP2001900	3.38	3.38	4.03	2.61	4.93	10.26		
NT2RP2001903	3.73	3.73	7.71	5.57	5.7	8.2		
NT2RP2001907	3.1	3.1	8.56	5.05	7.56	6.72		
NT2RP2001915	2.89	2.89	5.06	4.06	3.08	7.19		
NT2RP2001921	4.04	4.04	10.3	13.02	12.45	19.33	*	+
NT2RP2001926	2.75	2.75	8.25	3.55	5.3	5.64		
NT2RP2001933	5.65	5.65	52.55	43.62	43.55	48.58		
NT2RP2001936	1.54	1.54	5.03	2.8	2.96	3.8		
NT2RP2001943	25.33	25.33	49.4	47.71	40.48	51.65		
NT2RP2001946	3.05	3.05	4.3	3.41	4.51	6.1		
NT2RP2001947	3.18	3.18	3.44	3.93	3.21	6.88		
NT2RP2001948	3.59	3.59	10.79	5.71	7.29	19.72		
NT2RP2001956	5.24	5.24	12.73	11.54	9.42	9.89		
NT2RP2001969	4.05	4.05	7.82	3.24	5.7	6		
NT2RP2001976	2.9	2.9	6.39	5.68	6.95	6.41		
NT2RP2001978	3.26	3.26	6.08	4.18	4.83	6.03		
NT2RP2001985	2.14	2.14	3.8	2.56	4.63	2.51		
NT2RP2001991	3.34	3.34	5.7	1.53	4.62	5.19		
NT2RP2001997	3.16	3.16	8.43	5.31	7.47	6.98		
NT2RP2002015	136.21	136.21	265.98	266.8	340.89	272.43		
NT2RP2002017	3.24	3.24	6.06	2.06	3.3	2.57		
NT2RP2002025	6.08	6.08	51.73	31.83	26.94	37.84		
NT2RP2002030	6.06	6.06	11.95	9.77	8.07	8.76		
NT2RP2002032	2.31	2.31	4.95	2.39	3.81	2.55		
NT2RP2002033	3	3	6.71	3.53	7.57	8.05		
NT2RP2002041	3.5	3.5	6.37	2.76	3.15	9		
NT2RP2002046	3	3	6.88	2.99	7.4	6.01		
NT2RP2002047	2.71	2.71	3.86	2.06	3.87	3.83		
NT2RP2002050	7.67	7.67	11.66	9.78	6.47	7.77		
NT2RP2002052	3.77	3.77	8.39	6.6	3.99	6.28		
NT2RP2002058	2.07	2.07	4.11	3.1	4.82	3.39		
NT2RP2002060	1.48	1.48	3.44	2.1	5.9	3.18		
NT2RP2002063	3.61	3.61	5.83	3.3	5.2	6.65		
NT2RP2002066	10.11	10.11	13.47	5.47	9.88	9.73		
NT2RP2002070	3.74	3.74	6.3	2.15	3.1	3.83		
NT2RP2002076	3.72	3.72	5.1	3.35	5.58	3.63		

(1114)

特開 2002-191363

2225

2226

NT2RP2002078	13.09	13.09	105.74	73.87	88.51	76.08		
NT2RP2002079	5.76	5.76	36.34	31.87	32.66	36.54		
NT2RP2002099	4.19	4.19	5.82	4.7	6.75	6.03		
NT2RP2002105	2.66	2.66	12.25	11.41	14.04	11.23		
NT2RP2002115	1.63	1.63	5.21	1.98	3.42	1.52		
NT2RP2002124	3.66	3.66	6	4.56	5.38	3.88		
NT2RP2002137	3.99	3.99	4.83	2.21	2.1	1.76	**	-
NT2RP2002139	24.08	24.08	45.74	51.88	77.99	62.3	*	+
NT2RP2002154	1.37	1.37	4.13	3.56	2.36	3.2		
NT2RP2002155	351.63	351.63	869.83	623.53	501.61	620.68		
NT2RP2002172	1.5	1.5	3.33	2.78	3.53	5.13		
NT2RP2002185	3.29	3.29	7.65	7.3	8.56	8.12		
NT2RP2002188	1.74	1.74	5.95	4.15	4.31	4.95		
NT2RP2002192	2.9	2.9	7.6	6.65	6.42	5.83		
NT2RP2002193	5.21	5.21	5.22	4.76	4.95	5.75		
NT2RP2002208	5.96	5.96	7.31	4.7	5.67	5.14		
NT2RP2002219	2.2	2.2	1.8	2.22	3.2	2.26		
NT2RP2002231	1.72	1.72	3.1	4.11	2.76	5.76		
NT2RP2002232	2.59	2.59	5.17	3.93	4.7	6.08		
NT2RP2002235	5.62	5.62	15.07	16.26	16.18	15.18		
NT2RP2002239	37.02	37.02	67.99	72.09	67.21	63.77		
NT2RP2002252	2.64	2.64	3.66	2.63	2.76	2.94		
NT2RP2002256	4.62	4.62	15.3	11.37	16.99	12.91		
NT2RP2002257	7.01	7.01	22.77	18.65	25.09	20.6		
NT2RP2002259	1.58	1.58	13.91	9.9	12.15	10.49		
NT2RP2002264	0.6	0.6	3.14	3.2	3.12	3.92		
NT2RP2002267	3.66	3.66	8.75	8.95	8.3	11.16		
NT2RP2002270	4.26	4.26	8.23	16.09	10.47	14.71	*	+
NT2RP2002281	2.85	2.85	5.66	8.18	6.48	6.54	*	+
NT2RP2002288	4.32	4.32	6.6	5.33	5.56	4.23		
NT2RP2002292	5.42	5.42	8.4	6.64	8.08	6.95		
NT2RP2002299	9.6	9.6	9.22	11.75	19.42	15.97	*	+
NT2RP2002304	1.37	1.37	4.78	6.99	5.08	5.52		
NT2RP2002312	1.21	1.21	2.33	3.78	5.3	3.28	*	+
NT2RP2002316	3.28	3.28	5.43	7.57	7.21	8.2	**	+
NT2RP2002325	1.95	1.95	3.46	2.79	2.22	4.95		
NT2RP2002333	2.13	2.13	3.03	3.53	4.86	5.69	*	+
NT2RP2002371	5.43	5.43	9.14	9.72	12.07	11.65	*	+
NT2RP2002373	10.65	10.65	40.1	36.72	58.84	33.58		
NT2RP2002381	4.68	4.68	2.35	2.66	3.19	3.71		
NT2RP2002385	5.71	5.71	11.84	9.95	11.34	9.47		
NT2RP2002394	0.94	0.94	1.52	1.24	0.96	1.26		
NT2RP2002408	2.7	2.7	5.08	3.89	3.12	4.29		
NT2RP2002409	3.73	3.73	10.81	10.78	7.95	8.35		
NT2RP2002424	2.98	2.98	4.22	5.84	6.22	7.85	*	+
NT2RP2002426	6.44	6.44	11.38	7.59	8.46	8.93		
NT2RP2002429	17.2	17.2	24.73	27.87	33.96	20.83		
NT2RP2002437	4.61	4.61	5.98	4.83	6.47	4.79		
NT2RP2002439	3.83	3.83	6.69	2.68	3.22	4		
NT2RP2002442	13.63	13.63	71.65	57.78	63.05	78.84		

(1115)

特開2002-191363

2227

2228

NT2RP2002457	3.27	3.27	5.31	4.35	4.87	5.82		
NT2RP2002464	2.17	2.17	5.34	3.29	4.59	4.24		
NT2RP2002475	3.11	3.11	7.88	5.3	2.83	5.43		
NT2RP2002479	3.09	3.09	4.25	1.95	2.99	1.93		
NT2RP2002487	1.73	1.73	5.15	1.98	2.1	3.04		
NT2RP2002498	1.52	1.52	2.2	2.62	2.82	4.47		
NT2RP2002503	7.63	7.63	31.85	29.32	32.02	31.84		
NT2RP2002504	3.81	3.81	5.73	6	7.23	11.28		
NT2RP2002510	2.65	2.65	8.92	4.68	6.59	6.85		
NT2RP2002520	3.57	3.57	7.17	6.26	8.86	6.61		
NT2RP2002527	5.18	5.18	6.02	9	12.37	11.22	**	+
NT2RP2002533	3.34	3.34	6.27	4.83	6.94	5.88		
NT2RP2002537	3.22	3.22	4.02	4.09	5.91	10.08		
NT2RP2002542	4.81	4.81	4.64	5.99	5.93	9.73		
NT2RP2002546	4.31	4.31	5.85	6.5	4.91	5.24		
NT2RP2002549	4.06	4.06	9.33	7.68	10.49	11.65		
NT2RP2002564	4.11	4.11	11.18	10.67	9.21	9.29		
NT2RP2002591	2.45	2.45	7.03	3.31	4.79	5.79		
NT2RP2002595	9.67	9.67	12.41	12.06	13.39	14.79		
NT2RP2002602	4.19	4.19	7.53	5.68	8.96	10.42		
NT2RP2002606	1.27	1.27	2.93	2.26	2.97	3.95		
NT2RP2002609	6.12	6.12	9.95	4.79	5.48	7.74		
NT2RP2002618	2.74	2.74	6.83	4.2	6.34	5.44		
NT2RP2002621	4.24	4.24	10.22	6.58	7.52	9.79		
NT2RP2002643	1.79	1.79	4.84	3.11	5.98	3.94		
NT2RP2002672	4.48	4.48	9.23	8.03	9.37	9.87		
NT2RP2002673	4.13	4.13	5.01	8	12.88	17.73	*	+
NT2RP2002674	2.4	2.4	4.06	2.78	2.37	1.84		
NT2RP2002686	2.73	2.73	4.61	3.17	5.19	6.49		
NT2RP2002688	10.73	10.73	28.07	22.34	33.34	28.71		
NT2RP2002695	2.62	2.62	7.03	5.26	4.34	5.52		
NT2RP2002701	7.29	7.29	13.37	11.85	5.18	10.04		
NT2RP2002706	3.02	3.02	5.58	6.47	8.14	6.19	*	+
NT2RP2002710	11.2	11.2	36.97	39.43	33.9	42.75		
NT2RP2002721	5.53	5.53	9.42	7.33	7.34	8.45		
NT2RP2002727	3.56	3.56	6.87	2.17	3.96	3.52		
NT2RP2002734	3.59	3.59	6.65	5.71	7.65	7.54		
NT2RP2002736	5.25	5.25	13.13	13.17	13.59	17.58		
NT2RP2002740	2	2	5.11	3.18	2.81	3.13		
NT2RP2002741	2.12	2.12	4.8	4.68	5.39	5.78		
NT2RP2002750	2.5	2.5	9.22	7.73	9.57	8.18		
NT2RP2002752	4.59	4.59	10.39	7.93	7.66	8.03		
NT2RP2002753	4.49	4.49	9.91	8.66	9.04	14.67		
NT2RP2002760	4.79	4.79	10.31	3.56	4.5	4.13		
NT2RP2002769	4.42	4.42	6.06	3.89	5.67	6.43		
NT2RP2002778	4.13	4.13	10.59	6.8	8.2	9.12		
NT2RP2002791	8.89	8.89	54.27	48.75	53.08	50.19		
NT2RP2002800	1.66	1.66	4.52	4	5.19	5.38		
NT2RP2002805	3.38	3.38	5.46	4.75	6.44	3.81		
NT2RP2002811	3.27	3.27	8.23	5.87	8.13	10.61		

(1116)

特開2002-191363

2229

2230

NT2RP2002824	18.29	18.29	25.05	29.95	34.29	25.05		
NT2RP2002839	13.26	13.26	31.21	16.09	23.42	16.27		
NT2RP2002845	5.87	5.87	7.93	4.61	6.12	5.5		
NT2RP2002857	2.95	2.95	3.6	2.35	4.23	2.99		
NT2RP2002862	4.56	4.56	12.49	12.55	9.84	11.34		
NT2RP2002880	5.27	5.27	13.89	13.5	11.56	11.1		
NT2RP2002885	8.6	8.6	17.12	7.56	10.07	10.02		
NT2RP2002891	1.9	1.9	7.78	4.65	6.63	7.56		
NT2RP2002907	2.95	2.95	6.91	5.95	5.5	4.9		
NT2RP2002925	5.67	5.67	6.73	10.04	11.6	8.45	*	+
NT2RP2002927	10.08	10.08	10.63	19.05	21.85	15.48	**	+
NT2RP2002928	4.32	4.32	4.56	2.65	2.34	2.51	**	-
NT2RP2002929	3.96	3.96	9.74	8.09	7.86	9.87		
NT2RP2002934	1.5	1.5	1.4	2.81	3.01	2.49	**	+
NT2RP2002939	2.96	2.96	5.09	5.15	6.71	4.91		
NT2RP2002942	2.4	2.4	5.06	4.35	5.07	10.81		
NT2RP2002954	5.41	5.41	11.46	7.21	9.1	8.65		
NT2RP2002959	8.15	8.15	12.55	13.81	16.28	16.69	*	+
NT2RP2002974	5.03	5.03	6.53	4.7	3.45	4.54		
NT2RP2002976	6.92	6.92	17.08	11.84	14.66	12.42		
NT2RP2002979	4.41	4.41	8.12	7.03	8.66	7.6		
NT2RP2002980	6.44	6.44	15.09	15.56	11	17.45		
NT2RP2002986	3.87	3.87	7.6	6.68	7.4	7.39		
NT2RP2002987	3.52	3.52	8.23	11.1	9.18	9.4	*	+
NT2RP2002988	14.96	14.96	22.92	30.07	31.87	31.36	**	+
NT2RP2002993	2.97	2.97	4.18	3.8	3.84	2.84		
NT2RP2003000	4.88	4.88	8.34	6.97	9.62	9.97		
NT2RP2003008	4.85	4.85	5.06	3.34	4.76	4.78		
NT2RP2003020	4.45	4.45	44.26	28.35	46.52	34.33		
NT2RP2003032	1.91	1.91	4.02	5.82	6.48	6.59	**	+
NT2RP2003034	4.21	4.21	13.47	13.16	11.15	16.31		
NT2RP2003042	2.15	2.15	3.81	4.57	3.65	4.92		
NT2RP2003050	2.32	2.32	3.56	2.55	2.17	1.83		
NT2RP2003060	7.27	7.27	15.51	21.53	18.91	17.46	*	+
NT2RP2003073	5.61	5.61	8.73	7.06	10.51	8.17		
NT2RP2003099	5.05	5.05	3.67	3.21	3.73	2.84		
NT2RP2003108	3.6	3.6	4.23	5.29	3.91	6.62		
NT2RP2003115	1.68	1.68	5	7.75	4.69	4.84		
NT2RP2003117	2.71	2.71	5.69	3.6	4.66	4.13		
NT2RP2003121	1.83	1.83	3.47	4.03	2.69	3.33		
NT2RP2003125	4.13	4.13	11.44	15.42	12.55	13.66	*	+
NT2RP2003127	2.36	2.36	3.94	1.53	1.66	1.75		
NT2RP2003129	3.43	3.43	7.09	6.08	6.05	5.42		
NT2RP2003137	4.49	4.49	6.14	7.58	8.4	6.46	*	+
NT2RP2003138	4.66	4.66	20.24	16.55	17.45	16.92		
NT2RP2003146	6.2	6.2	24.78	18.5	23.25	25.96		
NT2RP2003148	3.09	3.09	6.73	3.06	4.6	4.04		
NT2RP2003150	1.45	1.45	5.71	3.98	5.2	4.3		
NT2RP2003157	6.93	6.93	34.27	34.29	31.85	32.84		
NT2RP2003158	6.3	6.3	25.32	26.87	28.69	59.31		

(1117)

特開2002-191363

2231

2232

NT2RP2003161	2.73	2.73	3.36	2.51	2.82	6.12		
NT2RP2003164	1.96	1.96	2.1	1.28	1.87	2.46		
NT2RP2003165	2.18	2.18	5.94	3.1	3.69	4.84		
NT2RP2003177	1.63	1.63	4.37	2.79	3.03	4.42		
NT2RP2003179	1.23	1.23	4.98	4.08	3.63	7.96		
NT2RP2003194	4.04	4.04	7.2	5.73	6.29	14.77		
NT2RP2003206	1.59	1.59	4.47	1.64	3.52	1.44		
NT2RP2003210	5.06	5.06	15.15	16.14	12.93	15.9		
NT2RP2003227	1.62	1.62	3.97	2.04	3.66	6.28		
NT2RP2003228	6.57	6.57	29.53	29.56	43.94	44.24		
NT2RP2003230	3.51	3.51	7.91	4.49	8.04	8.46		
NT2RP2003231	2.22	2.22	5.59	2.46	3.23	3.83		
NT2RP2003237	2.52	2.52	4.59	4.59	6.4	6.46	*	+
NT2RP2003239	2.3	2.3	4.46	2.97	4.46	4.05		
NT2RP2003243	2.16	2.16	4.13	2.38	3.28	3.98		
NT2RP2003265	3.93	3.93	5.33	4.22	4.88	4.92		
NT2RP2003267	2.73	2.73	3.15	3.24	4.17	7.42		
NT2RP2003272	6.03	6.03	14.8	16.93	23.85	32.58	*	+
NT2RP2003277	3.85	3.85	11.29	5.53	8.39	6.39		
NT2RP2003280	3.47	3.47	9.38	7.67	7.25	6.09		
NT2RP2003286	2.18	2.18	4.23	4.13	5	9.61		
NT2RP2003293	2.98	2.98	6.9	5.66	7.05	7.94		
NT2RP2003295	4.67	4.67	8.45	8.73	12.39	6.35		
NT2RP2003297	3.43	3.43	7.57	4.32	6.89	5.28		
NT2RP2003300	20.38	20.38	32.04	45.7	53.51	48.07	**	+
NT2RP2003302	2.88	2.88	4.52	3.46	3.81	7.26		
NT2RP2003307	0.57	0.57	2.62	1.43	1.49	1.38		
NT2RP2003308	1.44	1.44	4.5	2.77	4.44	5.44		
NT2RP2003311	4.18	4.18	5.83	7.35	4.25	8.2		
NT2RP2003329	2.99	2.99	4.41	2.63	3.89	4.44		
NT2RP2003339	3.06	3.06	7.01	3.76	4.92	3.64		
NT2RP2003345	4.15	4.15	8.38	2.77	3.97	6.33		
NT2RP2003347	2.55	2.55	4.23	2.08	1.98	3.46		
NT2RP2003367	2.15	2.15	4.65	2.7	1.98	1.44		
NT2RP2003369	1.34	1.34	4.71	2.16	2.36	0.89		
NT2RP2003383	4.05	4.05	6.75	7.66	7.17	6.99		
NT2RP2003390	9.1	9.1	17.93	16.66	14.27	12.94		
NT2RP2003391	9.39	9.39	12.9	11.96	9.91	12.84		
NT2RP2003393	4.23	4.23	6.99	6.14	5.03	9.44		
NT2RP2003394	8.67	8.67	16.21	17.56	21.75	17.23		
NT2RP2003401	4.39	4.39	5.97	3.52	3.72	2.9		
NT2RP2003403	3.42	3.42	7.64	6.62	8.55	7.79		
NT2RP2003433	3.02	3.02	15.54	13.62	13.5	13.08		
NT2RP2003445	3.2	3.2	4.74	4.08	3.5	3.93		
NT2RP2003446	2.67	2.67	6.23	5.06	6.05	4.47		
NT2RP2003456	2.04	2.04	6.57	4.26	5.89	3.81		
NT2RP2003466	3.56	3.56	20.09	17.34	25.96	23.53		
NT2RP2003469	6.2	6.2	5.65	6.19	7.53	6.33		
NT2RP2003470	5.64	5.64	6.47	5.06	6.11	6.44		
NT2RP2003471	2.72	2.72	3.88	3.22	2.78	3.38		

(1118)

特開2002-191363

2233

2234

NT2RP2003480	7.15	7.15	20.74	19.77	19.89	21.14		
NT2RP2003495	3.99	3.99	6.03	8.07	7.24	10.72	*	+
NT2RP2003499	1.52	1.52	4.58	3.67	4.05	2.75		
NT2RP2003505	0.98	0.98	3.21	2.62	3.88	1.4		
NT2RP2003506	2.54	2.54	6.53	5.65	5.36	4.78		
NT2RP2003511	3.67	3.67	5.57	4.22	3.1	2.96		
NT2RP2003513	3.79	3.79	6.01	5.49	5.57	5.71		
NT2RP2003517	2.9	2.9	2.52	1.32	1.11	0.85	**	-
NT2RP2003522	11.08	11.08	19.77	10.55	11.42	16.52		
NT2RP2003525	5.12	5.12	14.93	12.19	10.72	11.79		
NT2RP2003533	3.36	3.36	10.44	12.12	10.72	12.94		
NT2RP2003541	6.72	6.72	11.29	12.02	13.42	11.7		
NT2RP2003543	2.48	2.48	5.96	4.17	3.55	6.54		
NT2RP2003545	2.59	2.59	4.85	2.22	3.6	1.85		
NT2RP2003559	4.92	4.92	4.81	3.97	3.84	3.37	**	-
NT2RP2003564	4.46	4.46	3.93	2.53	1.97	2.42	**	-
NT2RP2003565	4.94	4.94	50.48	41.12	48.32	37.82		
NT2RP2003567	3.51	3.51	16.65	16.25	19.43	16.05		
NT2RP2003575	4.44	4.44	18.78	19.56	22.63	20.7		
NT2RP2003576	102.12	102.12	203.44	206.62	128.42	171.89		
NT2RP2003579	11.45	11.45	26.58	38.62	39.51	39.88	*	+
NT2RP2003581	3.85	3.85	6.1	4.33	4.38	3.96		
NT2RP2003587	8.37	8.37	11.47	13.35	14.11	12.14	*	+
NT2RP2003590	7.15	7.15	9.08	11.06	13.15	14.91	*	+
NT2RP2003593	1.58	1.58	4.57	7.84	4.43	8.59		
NT2RP2003596	4.86	4.86	10.86	14.43	13.12	17.96	*	+
NT2RP2003599	6.49	6.49	12.46	14.29	10.17	11.98		
NT2RP2003600	1.88	1.88	2.95	3.02	3.64	6.36		
NT2RP2003604	7.09	7.09	8.97	16.39	13.03	16.68	**	+
NT2RP2003629	3.72	3.72	5.25	3.11	4.56	2.38		
NT2RP2003630	4.09	4.09	6.66	4.79	6.78	3.84		
NT2RP2003643	5.49	5.49	4.88	7.15	9.8	8.51	*	+
NT2RP2003655	4.27	4.27	11.12	7.52	6.38	7.59		
NT2RP2003664	12.29	12.29	24.31	17.9	18.07	17.11		
NT2RP2003668	2.52	2.52	5.01	3.3	3.18	3.62		
NT2RP2003687	1.61	1.61	2.77	1.63	2.42	1.71		
NT2RP2003691	3.03	3.03	5.07	3.7	4.21	4.57		
NT2RP2003702	3.99	3.99	6.14	2.89	3.02	2.89		
NT2RP2003704	3.31	3.31	4.12	2.65	3.84	1.99		
NT2RP2003706	2.44	2.44	1.24	1.72	1.42	1.6		
NT2RP2003713	4.11	4.11	5.49	4.16	3.89	3.87		
NT2RP2003714	3.39	3.39	7.8	5.19	5.31	5.32		
NT2RP2003727	3.96	3.96	11.63	4.81	6.08	10.46		
NT2RP2003737	2.52	2.52	8.58	4.88	6.47	4.6		
NT2RP2003751	1.66	1.66	4.84	1.24	1.67	1.37		
NT2RP2003760	2.52	2.52	5.47	3.87	4.19	5.45		
NT2RP2003764	2.1	2.1	2.81	1.44	1.87	4.62		
NT2RP2003769	5.52	5.52	11.45	7.41	8.17	10.64		
NT2RP2003770	7.43	7.43	12.42	7.17	5.67	12.82		
NT2RP2003777	3.44	3.44	6.78	5.01	5.6	9.57		

(1119)

特開 2002-191363

2235

2236

NT2RP2003781	4.93	4.93	15.85	13.04	11.91	13.48		
NT2RP2003785	9.69	9.69	13.44	11.1	10.68	8.99		
NT2RP2003793	9.32	9.32	9.5	8.29	13.22	10.51		
NT2RP2003806	5.6	5.6	12.03	8.54	8.75	12.97		
NT2RP2003825	10.73	10.73	62.01	57.88	71.84	82.78		
NT2RP2003840	3.19	3.19	6.07	3.86	4.44	4.71		
NT2RP2003857	4.02	4.02	4.94	3.15	4.61	6.12		
NT2RP2003859	1.82	1.82	6	3.35	4.16	4.21		
NT2RP2003871	5.22	5.22	9.43	5.1	4.59	7.79		
NT2RP2003876	3.82	3.82	8.8	5.92	5.87	7.92		
NT2RP2003878	3.38	3.38	6.49	3.8	4.9	4.11		
NT2RP2003885	2.46	2.46	3.09	1.66	3.29	2.37		
NT2RP2003898	5.39	5.39	8.91	12.3	12.73	18.25	*	+
NT2RP2003902	5.09	5.09	10.78	8.24	8.23	10.42		
NT2RP2003912	3.83	3.83	14.48	5.91	7.43	6.74		
NT2RP2003931	1.81	1.81	6.03	3.95	6.86	4.42		
NT2RP2003940	2.31	2.31	9.51	7.1	6.2	7.3		
NT2RP2003950	2.81	2.81	5.48	3.84	5.57	2.98		
NT2RP2003952	1.86	1.86	5.63	2.58	4.23	2.98		
NT2RP2003968	4.82	4.82	7.38	9.86	11.76	13.51	*	+
NT2RP2003976	5.35	5.35	9.56	12.56	12.6	13.2	*	+
NT2RP2003981	3.27	3.27	7.41	4.62	2.03	4.07		
NT2RP2003984	5.57	5.57	15.87	10.21	4.25	10.34		
NT2RP2003986	2.79	2.79	6.22	6.29	6.32	5.17		
NT2RP2003988	2.36	2.36	6.84	4.51	7.42	5		
NT2RP2004013	8.46	8.46	13.75	14.68	13.19	17		
NT2RP2004014	4.24	4.24	10.07	4.06	5.12	4.08		
NT2RP2004036	6.88	6.88	14.85	14.08	19.02	16.03		
NT2RP2004041	2.77	2.77	5.02	3.96	4.43	5.19		
NT2RP2004042	1.99	1.99	4.6	4.41	2.02	4.09		
NT2RP2004049	4.68	4.68	19.13	14.24	15.5	16.3		
NT2RP2004060	5.7	5.7	10.41	7.09	8.67	10.84		
NT2RP2004066	2.17	2.17	4.31	3.05	4.83	3.65		
NT2RP2004069	3.99	3.99	7.24	3.54	6	4.26		
NT2RP2004076	3.73	3.73	5.82	1.92	4.2	3.61		
NT2RP2004080	4.21	4.21	9.26	4.45	6.47	6.15		
NT2RP2004081	3.27	3.27	5.39	3.51	3.71	4.5		
NT2RP2004098	2.32	2.32	6.48	5.4	3.1	5.75		
NT2RP2004108	3.82	3.82	9.56	7.18	5.89	7.56		
NT2RP2004124	3.13	3.13	5.9	3.68	5.82	3.92		
NT2RP2004130	3.67	3.67	9.32	5.51	9.12	8.4		
NT2RP2004133	2.05	2.05	6.41	3.69	6.54	6.25		
NT2RP2004141	5.72	5.72	7.15	5.14	7.05	7.05		
NT2RP2004142	5.33	5.33	8.1	4.18	5.45	3.93		
NT2RP2004152	3.34	3.34	4.78	5.7	7.49	4.39		
NT2RP2004165	3.71	3.71	8.3	5.87	5.92	6.54		
NT2RP2004170	1.86	1.86	5.97	5.37	4.17	4.94		
NT2RP2004172	2.93	2.93	5.24	4.69	5.58	4.26		
NT2RP2004176	3.45	3.45	8.4	7.77	10.21	8.98		
NT2RP2004179	4.01	4.01	9.17	3.94	5.07	4.15		

(1120)

特開 2002-191363

2237

2238

NT2RP2004187	3.16	3.16	6.36	3.87	3.88	4.59		
NT2RP2004190	5.1	5.1	5.46	5.49	7.33	9.98		
NT2RP2004194	7.54	7.54	14.57	18.5	23.44	19.83	*	+
NT2RP2004196	4.28	4.28	13.77	10.02	7.87	14.61		
NT2RP2004205	2.67	2.67	8.14	8.64	6.62	7.81		
NT2RP2004207	2.57	2.57	4.38	4.15	4.97	3.59		
NT2RP2004226	2.09	2.09	4.95	4.11	6.15	5.33		
NT2RP2004232	2.79	2.79	6.52	6	6.59	5.33		
NT2RP2004239	3.57	3.57	4.49	2.71	3.97	5.6		
NT2RP2004240	7.07	7.07	12.57	13	15.8	8.95		
NT2RP2004242	3.87	3.87	6.52	5.77	6.94	7.27		
NT2RP2004245	1.74	1.74	3.47	2.42	3.29	3.15		
NT2RP2004270	9.77	9.77	33.78	28.39	27.43	29.48		
NT2RP2004300	2	2	5.22	4.34	4.52	3.26		
NT2RP2004304	6.46	6.46	15.37	17.41	12.33	13.9		
NT2RP2004313	3.17	3.17	3.78	5.51	4.18	4.63	*	+
NT2RP2004316	3.46	3.46	5.84	4.9	4.96	4.04		
NT2RP2004321	4.71	4.71	6.06	6.79	7.43	6.29	*	+
NT2RP2004336	4.19	4.19	4.97	2.73	4.28	4.53		
NT2RP2004339	5.3	5.3	20.89	17.11	18.07	15.39		
NT2RP2004347	1.39	1.39	3.99	4.78	5	4.19		
NT2RP2004364	2.26	2.26	6.52	5.08	6.72	4.76		
NT2RP2004365	3.18	3.18	6.58	6.68	6.34	7.7		
NT2RP2004366	2.49	2.49	6.06	4.49	4.71	3.08		
NT2RP2004373	8.17	8.17	14.38	7.1	7.22	5.91		
NT2RP2004375	9.27	9.27	13.98	20.89	26.85	20.68	**	+
NT2RP2004389	5.25	5.25	5.62	5.01	6.26	5.61		
NT2RP2004392	8.88	8.88	23.7	13.04	19.48	20.89		
NT2RP2004396	1.98	1.98	6.27	6.65	4.98	6.17		
NT2RP2004399	5.24	5.24	8.12	12.56	7.74	9.52		
NT2RP2004400	2.07	2.07	3.55	2.36	3.47	2		
NT2RP2004404	15.79	15.79	46	45.56	40.89	41.74		
NT2RP2004410	16.64	16.64	24.04	27.99	33.46	32.69	*	+
NT2RP2004412	5.84	5.84	6.74	7.37	9.71	7.37		
NT2RP2004414	4.27	4.27	5.09	3.81	4.89	3.8		
NT2RP2004425	3.71	3.71	6.53	3.73	3.18	4.04		
NT2RP2004447	1.93	1.93	5.68	2.75	5.22	3.56		
NT2RP2004463	13.57	13.57	16.23	16.84	16.25	20.26		
NT2RP2004476	9.11	9.11	12.69	11.89	12.66	15.87		
NT2RP2004488	3.82	3.82	8.52	4.59	6.02	5.37		
NT2RP2004490	2.88	2.88	3.86	2.31	2.96	4.1		
NT2RP2004495	35.59	35.59	88.76	96.31	109.31	123.5	*	+
NT2RP2004512	4.25	4.25	7.62	5.84	6.41	7.12		
NT2RP2004523	5.18	5.18	11.04	8.44	7.56	10.21		
NT2RP2004524	2.19	2.19	6.33	4.97	5.32	5.56		
NT2RP2004536	8.99	8.99	16.96	14.1	14.51	17.51		
NT2RP2004538	8.03	8.03	24.44	20.15	25.59	22.24		
NT2RP2004548	4.45	4.45	9.92	7.39	9.1	10.51		
NT2RP2004551	4.95	4.95	5.62	7.17	9.98	6.21		
NT2RP2004556	83.73	83.73	210.17	226.48	298.92	241.84		

(1121)

特開 2002-191363

2239

2240

NT2RP2004568	5.19	5.19	11.18	6.52	9.16	9.99		
NT2RP2004580	3.98	3.98	7.71	5.71	7.88	7.04		
NT2RP2004585	11.28	11.28	49.82	36.69	46.91	64.56		
NT2RP2004587	1.85	1.85	4.16	2.07	2.89	3		
NT2RP2004594	4.56	4.56	9.24	11.47	12.21	28.18		
NT2RP2004600	3.49	3.49	5.76	2.22	3.22	3.09		
NT2RP2004602	4.62	4.62	6.32	6	8.49	6.26		
NT2RP2004606	392.21	392.21	581.19	612.4	897.5	764.63	*	+
NT2RP2004614	2.92	2.92	4.73	2.69	3.63	3.81		
NT2RP2004648	2.52	2.52	4.96	3.24	5.01	4.12		
NT2RP2004655	5.69	5.69	10.1	8.37	6.76	9.46		
NT2RP2004664	3.64	3.64	5.35	3.59	4.62	5.97		
NT2RP2004670	1.98	1.98	3.81	1.98	3.71	4.27		
NT2RP2004675	3.37	3.37	9.29	4.08	5.87	5.33		
NT2RP2004681	3.46	3.46	7.56	5.72	8.92	7.55		
NT2RP2004689	2.63	2.63	5.75	5.75	4.73	7.87		
NT2RP2004709	3.93	3.93	7.79	4.46	2.89	5.25		
NT2RP2004710	3.15	3.15	8.37	5.63	4.61	6.88		
NT2RP2004721	1.79	1.79	5.99	3.39	4.41	2.78		
NT2RP2004736	3.26	3.26	5.81	6.11	4.79	4.63		
NT2RP2004743	4.94	4.94	7.96	5.94	6.67	7.36		
NT2RP2004750	6.21	6.21	17.46	11.9	15.49	11.01		
NT2RP2004755	11.65	11.65	19.9	14.84	22.87	19.91		
NT2RP2004767	3.54	3.54	9	4.05	5.8	4.81		
NT2RP2004768	3.48	3.48	29.51	18.48	18.73	19.6		
NT2RP2004775	4.68	4.68	5.68	7.71	5.62	8.26		
NT2RP2004791	7.23	7.23	16.58	9.33	10.24	11.68		
NT2RP2004794	14.01	14.01	25.74	23.04	16.86	22.78		
NT2RP2004795	5.15	5.15	7.97	6.96	5.67	11.2		
NT2RP2004799	6.74	6.74	10.99	5.35	8.58	6.3		
NT2RP2004802	6.35	6.35	11.79	6.1	7.62	6.24		
NT2RP2004810	3.44	3.44	8.83	7.37	7.84	6.03		
NT2RP2004816	5.58	5.58	12.1	11.22	8.76	11.15		
NT2RP2004837	4.13	4.13	9.89	10.43	7.23	12.98		
NT2RP2004841	0.91	0.91	2.86	3.69	4.03	8.87		
NT2RP2004847	3.25	3.25	13.75	13.82	13.87	17.16		
NT2RP2004861	2.3	2.3	5.23	2.33	4.23	2.46		
NT2RP2004897	3.35	3.35	6.43	4.26	3.27	3.35		
NT2RP2004932	6.64	6.64	10.16	7.96	8.53	6.91		
NT2RP2004933	4.63	4.63	3.41	2.98	2.93	3.2	*	-
NT2RP2004936	3.69	3.69	6.41	4.56	4.42	7.53		
NT2RP2004951	2.98	2.98	10.48	5.09	5.22	19.28		
NT2RP2004959	3.13	3.13	6.61	6.43	6.26	6.5		
NT2RP2004961	2.1	2.1	4.79	4.89	6.49	5.44		
NT2RP2004962	2.27	2.27	7.28	4.5	5.57	4.47		
NT2RP2004966	2.26	2.26	6.07	4.1	4.1	2.97		
NT2RP2004967	3.87	3.87	6.16	4.07	4.82	3.26		
NT2RP2004974	5.27	5.27	5.43	3.59	3.47	3.89	**	-
NT2RP2004978	2.68	2.68	5.26	4.17	6.39	5.09		
NT2RP2004982	0.57	0.57	1.82	2.2	1.94	1.72		

(1122)

特開 2002-191363

2241

2242

NT2RP2004985	16.03	16.03	45.34	44.65	46.12	54.4		
NT2RP2004999	2.21	2.21	5.64	4.27	8.86	10.34		
NT2RP2005000	3.62	3.62	5.76	4.33	4.76	4.65		
NT2RP2005001	5.41	5.41	7.91	8.26	9.15	8.32		
NT2RP2005003	3.8	3.8	7.2	6.11	7.91	6.2		
NT2RP2005012	6.61	6.61	20.14	18.41	20.96	17.87		
NT2RP2005018	1.9	1.9	4.24	3.29	2.24	2.91		
NT2RP2005020	6.12	6.12	23.58	19.97	19.94	22.96		
NT2RP2005022	1.65	1.65	5.01	5.09	7.24	4.77		
NT2RP2005027	5.96	5.96	38.61	42.51	40.08	33.77		
NT2RP2005031	1.54	1.54	4.99	3.94	4.53	3.92		
NT2RP2005035	44.19	44.19	94.82	116.52	107.36	106.69	*	+
NT2RP2005037	4.28	4.28	5.87	7.91	10.26	7.09	*	+
NT2RP2005038	4.86	4.86	4.84	1.85	2.29	3.1	**	-
NT2RP2005048	9.92	9.92	30.91	30.14	33.38	32.48		
NT2RP2005069	16.01	16.01	34.88	21.99	20.63	27.64		
NT2RP2005073	7.36	7.36	30.35	29.36	28.24	30.33		
NT2RP2005097	2.39	2.39	5.6	5.27	5.2	4.6		
NT2RP2005108	1.76	1.76	3.95	2.84	4.21	7.12		
NT2RP2005116	3.53	3.53	5.96	6.27	5.42	5.89		
NT2RP2005126	5.88	5.88	8.31	8.9	14.96	8.51		
NT2RP2005135	5.08	5.08	5.22	4.65	6.59	5.47		
NT2RP2005139	1.94	1.94	2.77	1.87	1.81	2.45		
NT2RP2005140	3.82	3.82	4.86	12.39	6.72	8.55	*	+
NT2RP2005144	4.04	4.04	6.31	6.69	5.97	9.7		
NT2RP2005147	2.23	2.23	5.49	5.61	6.15	6.25		
NT2RP2005148	2.86	2.86	5.63	3.83	6.65	4.83		
NT2RP2005159	3.92	3.92	5.6	4.94	6.38	7.41		
NT2RP2005162	3.23	3.23	5.56	4.57	5.4	4.21		
NT2RP2005163	9.15	9.15	20.61	24.53	28.92	23.77	*	+
NT2RP2005168	2.87	2.87	6.14	5.24	4.79	4.88		
NT2RP2005181	2.64	2.64	5.42	3.4	2.11	1.98		
NT2RP2005204	5.4	5.4	7.81	9.08	11.94	11.81	*	+
NT2RP2005219	4.61	4.61	9.64	7.09	10.28	8.7		
NT2RP2005227	3.59	3.59	10.43	7.55	5.36	9.97		
NT2RP2005237	26.49	26.49	94.81	86.96	105.8	93.92		
NT2RP2005239	2.24	2.24	6.07	2.62	4.27	4.34		
NT2RP2005247	10.63	10.63	37.59	35.58	46.1	46.95		
NT2RP2005254	4.35	4.35	9.14	5.7	6.44	6.93		
NT2RP2005270	9.06	9.06	17.44	10.82	9.28	17.11		
NT2RP2005276	7.19	7.19	11.53	10.88	11.68	15.71		
NT2RP2005287	7.98	7.98	11.97	8.37	7.7	13.36		
NT2RP2005288	2.51	2.51	5.14	2.89	5.59	5.22		
NT2RP2005289	4.26	4.26	8.48	6.68	9.08	7.49		
NT2RP2005293	5	5	6.93	13.68	14.37	15.66	**	+
NT2RP2005315	5.79	5.79	10.64	8.04	12.95	16.68		
NT2RP2005322	5.05	5.05	15.42	18.91	11.33	22.43		
NT2RP2005325	8.45	8.45	18.4	15.57	13.63	20.01		
NT2RP2005336	1.71	1.71	6.68	4.18	5.74	5.3		
NT2RP2005343	2.44	2.44	7.48	3.91	4.11	5.89		

(1123)

特開2002-191363

2243

2244

NT2RP2005344	3.39	3.39	4.83	2.37	2.67	3.32		
NT2RP2005347	3.14	3.14	3.61	3.34	2.96	3.53		
NT2RP2005354	6.49	6.49	11.79	10.37	13	11.38		
NT2RP2005358	35.87	35.87	109.04	101.37	134.72	117.96		
NT2RP2005360	2.93	2.93	5	3.59	4.97	3.84		
NT2RP2005378	5.27	5.27	13.12	7.54	8.7	13.93		
NT2RP2005391	3.06	3.06	5.41	4.21	6.76	7.72		
NT2RP2005393	1.61	1.61	6.34	4.86	6.16	4.07		
NT2RP2005407	2.59	2.59	5.71	4.28	5.65	4.64		
NT2RP2005419	2.65	2.65	9.05	6.37	8.5	6.77		
NT2RP2005425	5.63	5.63	18.38	15.27	18.89	15.46		
NT2RP2005429	3.23	3.23	5.85	5.41	6.65	5.64		
NT2RP2005436	4.65	4.65	10.5	7.02	4.28	4.97		
NT2RP2005441	2.28	2.28	5.62	3.36	3.77	5.79		
NT2RP2005442	24.92	24.92	40.66	34.62	25.56	41.66		
NT2RP2005444	10.72	10.72	19.24	21.92	21.07	25.56	*	+
NT2RP2005453	2.79	2.79	7.44	2.63	4.09	3.15		
NT2RP2005457	15.12	15.12	23.21	28.69	37.38	31.61	*	+
NT2RP2005458	2.47	2.47	5.27	3.55	4.16	4.95		
NT2RP2005463	7.73	7.73	15.23	15.65	22.11	25.05	*	+
NT2RP2005464	5.96	5.96	11.91	9.22	4.67	10.35		
NT2RP2005465	1.81	1.81	6.69	3.86	3.75	3.74		
NT2RP2005472	10.98	10.98	32.59	28.21	27.9	25.85		
NT2RP2005476	5.01	5.01	8.99	7.01	6.98	6.08		
NT2RP2005490	7.51	7.51	21.09	18.18	25.55	23.45		
NT2RP2005491	4.99	4.99	12.47	8.63	10.12	8.78		
NT2RP2005495	3.56	3.56	5.77	3.38	4.55	4.3		
NT2RP2005496	4.84	4.84	18.25	11.3	13.16	11.28		
NT2RP2005498	2.92	2.92	7.45	5.18	5.03	4.98		
NT2RP2005501	2.04	2.04	5.54	3.12	4.34	2.46		
NT2RP2005506	124.3	124.3	217.82	139.27	121.83	104.81		
NT2RP2005509	6.97	6.97	10.45	11.4	9.61	15.73		
NT2RP2005514	3.93	3.93	6	4.06	7.05	4.39		
NT2RP2005520	14.95	14.95	32.39	27.11	39.97	33.03		
NT2RP2005525	6.19	6.19	7.01	7.81	7.68	4.79		
NT2RP2005531	2.18	2.18	3.33	1.67	2.12	1.9		
NT2RP2005535	4.66	4.66	9.09	9.34	7.79	8.91		
NT2RP2005539	3.39	3.39	6.22	6.43	5.84	7.45		
NT2RP2005540	3.2	3.2	7.15	4.79	5.58	6.59		
NT2RP2005541	21.25	21.25	39.57	25.85	38.31	39.61		
NT2RP2005549	2.69	2.69	7.66	6.72	4.85	7.11		
NT2RP2005555	7.97	7.97	10.1	14.96	16.19	15.37	**	+
NT2RP2005557	4.89	4.89	8.47	4.03	6.52	6.26		
NT2RP2005581	3.93	3.93	9.61	6.32	7.95	6.89		
NT2RP2005586	1.56	1.56	3.18	3.21	2.92	4.74		
NT2RP2005597	2.77	2.77	2.93	2.98	4.1	3.84		
NT2RP2005600	1.81	1.81	3.71	4.03	4.29	4.44	*	+
NT2RP2005605	4.93	4.93	14.29	13.17	15.14	15.75		
NT2RP2005614	3.06	3.06	5.62	3.68	4.11	2.45		
NT2RP2005620	3.47	3.47	6.26	3.6	3.92	3.11		

(1124)

特開 2002-191363

2245

2246

NT2RP2005622	6.14	6.14	5.07	6.21	7.43	4.61		
NT2RP2005632	5.72	5.72	10.95	11.57	10.42	14.89		
NT2RP2005635	2.22	2.22	19.06	18.14	23.77	18.14		
NT2RP2005637	1.53	1.53	8	3.73	3.71	4.14		
NT2RP2005640	1.72	1.72	7.22	7.49	8.73	6.06		
NT2RP2005645	4.68	4.68	11.8	10.61	11.47	9.67		
NT2RP2005651	3.45	3.45	7.88	7.64	6.78	10.15		
NT2RP2005654	4.08	4.08	4.14	3.02	2.52	3.8		
NT2RP2005666	4.91	4.91	5.27	4.34	7.7	4.74		
NT2RP2005669	7.15	7.15	7.95	7.05	11.14	8.21		
NT2RP2005670	2.35	2.35	6.91	7.77	5.04	5.2		
NT2RP2005671	3.12	3.12	7.83	10.77	8.9	9.78	*	+
NT2RP2005675	7.32	7.32	37.84	34.46	40.94	40.02		
NT2RP2005683	2.56	2.56	7.01	7.16	5.19	7.16		
NT2RP2005690	2.84	2.84	4.48	2.82	3.74	3.4		
NT2RP2005694	4.07	4.07	5.49	3.77	6.26	3.54		
NT2RP2005701	5.97	5.97	8.82	10.39	10.35	9.52	*	+
NT2RP2005712	5.67	5.67	5.28	4.83	7.94	6.33		
NT2RP2005719	1.86	1.86	3.26	4.42	3.8	3.76	*	+
NT2RP2005722	4.16	4.16	11.13	13.39	15.7	15.94	*	+
NT2RP2005723	2.71	2.71	4.2	3.65	4.58	3.67		
NT2RP2005726	2.55	2.55	4.13	2.86	4.01	3.22		
NT2RP2005729	4.64	4.64	9.94	10.21	10.7	10.62		
NT2RP2005731	3.05	3.05	3.39	2.51	2.16	1.27	*	-
NT2RP2005732	9.41	9.41	57.73	48.37	75.21	41.64		
NT2RP2005737	10.75	10.75	22.28	27.16	25.02	17.59		
NT2RP2005741	3.03	3.03	5.35	3.68	3.31	3.37		
NT2RP2005748	1.86	1.86	5.94	3.8	3.72	2.95		
NT2RP2005752	2.46	2.46	5.55	3.27	4.37	3.8		
NT2RP2005753	8.45	8.45	14.76	11.12	11.69	14.45		
NT2RP2005763	3	3	8.03	4.22	4.77	5		
NT2RP2005767	3.72	3.72	7.79	5.55	6.75	6.29		
NT2RP2005773	8.11	8.11	10.02	10.6	11.59	13.37	*	+
NT2RP2005774	4.25	4.25	12.72	6.86	12.24	14.61		
NT2RP2005775	3.75	3.75	7.2	3.35	4.83	5.63		
NT2RP2005781	5.11	5.11	9.88	9.82	7.19	12.94		
NT2RP2005784	5.41	5.41	11.51	7.68	12.12	14.06		
NT2RP2005789	3.98	3.98	11.24	7.89	9.52	8.86		
NT2RP2005799	2.45	2.45	6.35	2.64	4.67	3.7		
NT2RP2005804	9.01	9.01	25	27.85	27.32	30.57		
NT2RP2005812	2.63	2.63	4.83	2.9	3.89	5.21		
NT2RP2005815	2.48	2.48	3.15	2.38	3.21	5.2		
NT2RP2005835	5.99	5.99	11.26	7.13	13.74	11.69		
NT2RP2005841	2.32	2.32	10.04	4.89	7.43	11.23		
NT2RP2005853	1.29	1.29	4.44	2.71	4.6	4.96		
NT2RP2005857	7.37	7.37	9.87	13.46	7.93	20.27		
NT2RP2005859	2.76	2.76	5	2.91	6.14	4.78		
NT2RP2005860	1.41	1.41	3.54	1.45	1.89	2.22		
NT2RP2005863	3.03	3.03	6.55	10.76	18.29	15.94	*	+
NT2RP2005868	3.86	3.86	5.85	5.1	6.3	7.77		

(1125)

特開 2002-191363

2247

2248

NT2RP2005876	5.7	5.7	12.31	7.84	8.29	8.2		
NT2RP2005878	2.26	2.26	8.44	5.25	4.95	6.32		
NT2RP2005883	13.54	13.54	21.06	23.75	9.57	28.09		
NT2RP2005886	7.18	7.18	50.05	51.13	62.09	50.14		
NT2RP2005887	3.76	3.76	6.51	4.74	8.05	4.81		
NT2RP2005890	4.17	4.17	9.77	11.87	17.13	12.15	*	+
NT2RP2005901	3.19	3.19	5.69	3.91	6.18	5.14		
NT2RP2005902	3.17	3.17	4.33	4.78	4.77	6.25		
NT2RP2005908	3.09	3.09	7.86	4.89	3.7	6.34		
NT2RP2005927	1.77	1.77	2.66	2.25	4.36	3.07		
NT2RP2005933	2.5	2.5	5.59	6.4	4.77	6.42		
NT2RP2005941	2.09	2.09	5.2	3.31	4.41	3.9		
NT2RP2005942	4	4	6.86	3.08	4.59	4.64		
NT2RP2005946	4.63	4.63	9.49	5.33	7.06	6.24		
NT2RP2005970	5.44	5.44	14	16.16	22.05	18.9	*	+
NT2RP2005980	3.71	3.71	5.25	2.69	3.46	2.37		
NT2RP2005994	2.99	2.99	6.76	4.28	3.28	5.14		
NT2RP2006004	1.31	1.31	2.89	2.07	6.09	2.58		
NT2RP2006013	1.38	1.38	4.91	3.1	5.07	4.92		
NT2RP2006023	8.37	8.37	17.77	20	21.43	21.14	*	+
NT2RP2006028	5.03	5.03	10.23	7.47	9.89	9.71		
NT2RP2006038	4.67	4.67	5.86	2.79	5.4	1.09		
NT2RP2006042	8.3	8.3	7.22	6.63	5.89	6.3	*	-
NT2RP2006043	5.65	5.65	7.59	7.6	10.99	8.29		
NT2RP2006052	1.48	1.48	4.48	4.13	3.12	4.54		
NT2RP2006057	3.73	3.73	6.23	5.69	3.83	4.95		
NT2RP2006064	4.16	4.16	7.73	5.86	6.81	9.08		
NT2RP2006068	2.76	2.76	6.75	6.8	7.81	5.81		
NT2RP2006069	1.46	1.46	4.94	3.56	3.95	3.3		
NT2RP2006071	8.37	8.37	7.8	9.28	10.48	9.11	*	+
NT2RP2006090	6.62	6.62	5.78	3.27	3.55	3.64	**	-
NT2RP2006092	3.78	3.78	8.3	6.18	8.04	7.07		
NT2RP2006097	14.05	14.05	40.38	31.2	25.81	40.02		
NT2RP2006098	1.94	1.94	4.27	4.52	4.61	7.65		
NT2RP2006099	3.84	3.84	11.02	10.65	10.99	13.34		
NT2RP2006100	2.87	2.87	5.78	3.63	7.31	5.19		
NT2RP2006103	2.39	2.39	5.54	2.6	3.93	1.71		
NT2RP2006106	6.48	6.48	21.51	18.05	24.81	22.3		
NT2RP2006127	3.17	3.17	4.92	1.62	1.26	1.21	*	-
NT2RP2006134	4.25	4.25	4.41	6.08	6.7	5.47	**	+
NT2RP2006141	3.91	3.91	7.94	7.45	6.04	9.08		
NT2RP2006166	3.1	3.1	10.65	9.01	8.94	7.85		
NT2RP2006176	2.15	2.15	4.26	3.95	5.73	4.69		
NT2RP2006181	1.68	1.68	2.84	3.21	3.14	2.45		
NT2RP2006184	8.85	8.85	17.16	20.8	19.95	17.1		
NT2RP2006186	3.01	3.01	4.57	2.77	2.29	4.33		
NT2RP2006196	5.24	5.24	7.21	5.25	5.23	4.16		
NT2RP2006199	5.06	5.06	4.38	3.81	3.65	3.64	**	-
NT2RP2006200	0.87	0.87	3.43	4.37	4.52	2.17		
NT2RP2006210	20.08	20.08	59.85	75.37	70.55	96.59	*	+

(1126)

特開 2002-191363

2249

2250

NT2RP2006219	2.88	2.88	6.26	5.97	5.11	7.36
NT2RP2006224	3.7	3.7	7.55	9	7.7	8.93
NT2RP2006237	1.97	1.97	4.79	3.45	2.74	4.14
NT2RP2006238	3.9	3.9	6.33	4.2	4.69	3.93
NT2RP2006258	4.5	4.5	6.73	3.07	4.27	4.39
NT2RP2006261	7.32	7.32	3.98	2.04	3.19	7.69
NT2RP2006269	4.11	4.11	7.96	9.52	5.46	9.06
NT2RP2006275	3.67	3.67	30.36	23.46	35.36	25.14
NT2RP2006282	3.16	3.16	8.89	8.85	8.4	7.05
NT2RP2006302	5.69	5.69	12.68	13.12	12.4	11.87
NT2RP2006312	4.88	4.88	8.22	8.47	9.13	9.8
NT2RP2006320	4.27	4.27	9.87	6.42	9.32	9.69
NT2RP2006321	3.27	3.27	4.23	2.79	4.99	4.13
NT2RP2006323	4.1	4.1	2.59	2.39	3.6	1.83
NT2RP2006333	0.67	0.67	1.82	1.7	1.04	1.17
NT2RP2006334	2.24	2.24	4.02	4.57	2.72	3.54
NT2RP2006338	2.4	2.4	5.26	4.73	5.11	4.04
NT2RP2006339	2.24	2.24	2.94	2.47	1.93	2.06
NT2RP2006355	3.61	3.61	4.59	3.14	3.39	2.22
NT2RP2006365	3.3	3.3	4.44	2.42	2.6	1.3
NT2RP2006374	16.34	16.34	111.62	108.73	174.7	73.65
NT2RP2006393	4.93	4.93	7.68	7.38	8	6.95
NT2RP2006394	8.59	8.59	17.91	11.3	11.18	15.38
NT2RP2006400	2.25	2.25	4.51	2.08	3.58	1.95
NT2RP2006411	27.71	27.71	42.11	23.61	17.25	37.31
NT2RP2006429	2.22	2.22	7.3	2.82	5.3	2.21
NT2RP2006435	1.46	1.46	5.29	1.76	2.65	1.98
NT2RP2006436	2.33	2.33	6.43	4.33	5.28	3.75
NT2RP2006441	4.69	4.69	8.19	7.76	8.89	8.37
NT2RP2006447	2.41	2.41	4.78	3.18	2.63	3.87
NT2RP2006454	2.58	2.58	5.38	4.39	3.37	4.03
NT2RP2006455	3.79	3.79	7.14	2.91	4.62	9.23
NT2RP2006456	1.96	1.96	5.99	2.51	4.49	3.17
NT2RP2006464	5.44	5.44	8.28	4.47	8.85	7.9
NT2RP2006467	4.17	4.17	10	8.56	12.47	12.58
NT2RP2006472	5.05	5.05	6.84	7.24	6.92	7.37
NT2RP2006474	4.69	4.69	16.3	18.19	32.31	21.3
NT2RP2006475	2.5	2.5	9.54	6.14	6.86	7.66
NT2RP2006476	5.34	5.34	14.94	7.62	13.82	17.24
NT2RP2006501	2.44	2.44	7.28	4.6	7.45	7.74
NT2RP2006512	10.25	10.25	19.79	16.72	7.89	29.01
NT2RP2006526	2.09	2.09	5.19	2.24	2.78	2.31
NT2RP2006527	3.61	3.61	7.05	4.56	6.14	6.46
NT2RP2006534	2.24	2.24	4.49	2.08	2.95	2.73
NT2RP2006537	6.08	6.08	15.7	11.72	17.73	12.82
NT2RP2006543	7.83	7.83	14.8	6.52	5.4	6.88
NT2RP2006554	1.33	1.33	3.71	1.79	3.76	2.2
NT2RP2006565	3.78	3.78	8.91	5.79	8.42	7.55
NT2RP2006571	1.38	1.38	3.88	2.77	4.01	2.29
NT2RP2006573	2.1	2.1	4.02	3.05	3.6	2.41

(1127)

特開2002-191363

2251

2252

NT2RP2006598	2.25	2.25	7.04	4.34	6.56	4.78		
NT2RP2006601	24.92	24.92	35.13	38.45	45.47	31.69		
NT2RP3000002	5.04	5.04	6.09	4.7	5.04	8.18		
NT2RP3000011	1.82	1.82	5.9	2.59	1.85	2.22		
NT2RP3000014	3.29	3.29	7.66	4.22	3.06	4.95		
NT2RP3000016	3.42	3.42	7	5.29	6.56	6.11		
NT2RP3000022	1.71	1.71	3.93	1.72	4.68	0.59		
NT2RP3000024	3.74	3.74	7.03	4.31	4.92	4.06		
NT2RP3000031	4.66	4.66	8.66	4	6.75	4.86		
NT2RP3000034	3.76	3.76	6.24	4.44	7.13	3.23		
NT2RP3000037	2.76	2.76	6.5	9.41	13.44	11.06	*	+
NT2RP3000040	2.04	2.04	5.96	3.21	3.46	3.56		
NT2RP3000041	2.15	2.15	7.35	3.71	3.01	3.16		
NT2RP3000046	1.95	1.95	4.42	3.67	7.11	3.84		
NT2RP3000047	3.25	3.25	5.55	5.85	6.2	5.94		
NT2RP3000049	2.54	2.54	6.94	5.26	7.78	3.58		
NT2RP3000050	4.99	4.99	9.03	3.76	8.5	6.22		
NT2RP3000051	5.99	5.99	10.69	8.51	11.19	9.72		
NT2RP3000054	4.31	4.31	6.5	4.38	5.35	3.22		
NT2RP3000055	1.98	1.98	4.76	3.81	2.67	3.96		
NT2RP3000056	2.87	2.87	7.09	5.59	3.32	3.91		
NT2RP3000059	2.54	2.54	5.1	1.89	4.07	1.6		
NT2RP3000063	2.18	2.18	5.51	3.34	5.19	2.27		
NT2RP3000068	3.76	3.76	24.22	25.83	37.88	23.13		
NT2RP3000069	17.44	17.44	20.58	22	28.87	18.2		
NT2RP3000072	5.9	5.9	6.18	4.96	5.39	4.19	*	-
NT2RP3000080	4.38	4.38	6.72	3.78	5.28	3.93		
NT2RP3000085	1.9	1.9	4.84	5.13	4.66	5.5		
NT2RP3000087	3.77	3.77	9.1	6.22	5.61	6		
NT2RP3000092	1.92	1.92	3.6	2.72	3.2	2.52		
NT2RP3000109	1.74	1.74	5.05	5.63	7.94	4.24		
NT2RP3000119	4.66	4.66	14.27	11.29	13.7	14.28		
NT2RP3000125	3.02	3.02	5.56	3.42	4.53	2		
NT2RP3000131	7.84	7.84	14.37	16.23	19.96	12.93		
NT2RP3000134	5.96	5.96	9.01	6.61	7.25	6.46		
NT2RP3000137	3.88	3.88	6.48	5.58	6.3	6.11		
NT2RP3000142	2.87	2.87	7.77	7.28	5.03	5.31		
NT2RP3000148	1.84	1.84	6.28	4.9	5.04	5.34		
NT2RP3000149	2.51	2.51	6.97	6.14	7.77	8.24		
NT2RP3000163	2.16	2.16	6.17	3.27	3.9	2.5		
NT2RP3000168	5.53	5.53	14.55	12.8	11.65	11.73		
NT2RP3000169	3.74	3.74	6.01	6.03	8.47	5.72		
NT2RP3000171	10.86	10.86	16.71	28.33	38.98	25.93	*	+
NT2RP3000172	0.86	0.86	1.53	1.66	1.2	1.46		
NT2RP3000186	4.32	4.32	10.6	19.18	15.43	15.82	*	+
NT2RP3000197	1.22	1.22	3.66	4.03	4.29	3.39		
NT2RP3000201	2.4	2.4	7.2	10.49	8.4	7.88		
NT2RP3000204	2.16	2.16	4.44	3.88	4.1	4.25		
NT2RP3000207	2.87	2.87	4.71	3	2.6	2.45		
NT2RP3000216	5.38	5.38	10.1	5.87	9.5	5.73		

(1128)

特開2002-191363

2253

2254

NT2RP3000220	5.14	5.14	5.66	3.68	5.69	2.92		
NT2RP3000221	2.18	2.18	5.45	6.26	6.63	5.93		
NT2RP3000232	2.7	2.7	8.01	7.1	5.52	5.92		
NT2RP3000233	1.55	1.55	6.01	6.9	5.91	4.06		
NT2RP3000234	3.23	3.23	9.09	12.89	10.4	11.41	*	+
NT2RP3000235	1.57	1.57	3.3	2.35	2.92	1.38		
NT2RP3000239	4.61	4.61	11.11	9.51	9.71	14.92		
NT2RP3000247	3.25	3.25	5.82	2.92	4.04	1.96		
NT2RP3000251	6.11	6.11	6.52	5.22	5.82	3.25		
NT2RP3000252	3.73	3.73	7.99	7.61	8.53	8.4		
NT2RP3000255	2.18	2.18	2.96	3.26	3.13	1.97		
NT2RP3000262	6.72	6.72	9.43	11.67	7.95	9.13		
NT2RP3000266	6.47	6.47	15.5	13.38	10.83	12.64		
NT2RP3000267	2.71	2.71	4.04	2.9	2.64	3.03		
NT2RP3000271	4.38	4.38	5.57	5.11	4.84	3.72		
NT2RP3000278	7.84	7.84	56.85	48.55	82.07	42.57		
NT2RP3000281	4.94	4.94	10.72	8.19	8.22	7.27		
NT2RP3000292	5.63	5.63	14.1	9.17	6.77	6.93		
NT2RP3000299	2.31	2.31	4.92	3.73	4.89	4.98		
NT2RP3000304	2.15	2.15	3.48	2.85	3.36	1.64		
NT2RP3000310	7.24	7.24	24.22	18.94	23.07	19.88		
NT2RP3000312	2.99	2.99	8.16	3.31	5.25	3.87		
NT2RP3000320	7.06	7.06	6.17	5.25	4.74	4.74	**	-
NT2RP3000322	11.05	11.05	18.76	32.59	45.13	46.95	**	+
NT2RP3000324	6.91	6.91	46.42	36.64	43.53	39.68		
NT2RP3000326	1.95	1.95	6.17	4.02	5.75	3.53		
NT2RP3000329	2.5	2.5	5.96	4.97	8.84	5.9		
NT2RP3000330	4.1	4.1	6.18	4.62	5.53	6.12		
NT2RP3000333	3.23	3.23	7.45	4.36	5.28	4.52		
NT2RP3000341	8.8	8.8	12.85	14.81	18.59	14.41	*	+
NT2RP3000344	2.73	2.73	3.75	2.69	3.54	2.29		
NT2RP3000345	3.09	3.09	3.57	1.65	1.97	2.66	*	-
NT2RP3000348	444.59	444.59	802.63	824.62	1016.01	909.68		
NT2RP3000350	4.25	4.25	10.34	4.57	9.28	6.4		
NT2RP3000359	9.53	9.53	24.44	8.54	11.36	16.62		
NT2RP3000361	7.5	7.5	11.12	7.89	7.81	8.95		
NT2RP3000366	7.38	7.38	14.27	9.52	11.84	16.13		
NT2RP3000378	2.67	2.67	5.75	3.92	4.78	2.47		
NT2RP3000384	5.42	5.42	10.88	9.52	13.1	9.28		
NT2RP3000389	12.54	12.54	21.49	23.95	35.02	27.32	*	+
NT2RP3000393	3.74	3.74	6.16	5.03	4.53	4.77		
NT2RP3000395	110.27	110.27	212	108.33	38.18	148.45		
NT2RP3000397	2.83	2.83	5.28	2.51	5.26	3.31		
NT2RP3000398	3.39	3.39	10.12	11.46	11.18	12.26		
NT2RP3000403	3.22	3.22	9.39	10.1	8.2	8.44		
NT2RP3000418	3.4	3.4	10.22	7.12	11.08	13.42		
NT2RP3000424	2.86	2.86	9.43	6.25	9.52	6.86		
NT2RP3000427	4.65	4.65	9.05	11.55	13.43	12.35	*	+
NT2RP3000431	2.05	2.05	4.93	3.43	3.26	3.93		
NT2RP3000433	2.63	2.63	8.65	5.65	7.09	6.65		

(1129)

特開2002-191363

2255

2256

NT2RP3000436	11.39	11.39	20.93	18.76	9.35	18.86		
NT2RP3000439	1.4	1.4	3.61	2.54	3.56	2		
NT2RP3000441	3.88	3.88	7.4	7.56	7.92	6.39		
NT2RP3000444	3.31	3.31	7.29	2.36	3.25	2.2		
NT2RP3000448	4.45	4.45	10.15	4.05	6.54	3.93		
NT2RP3000449	2.84	2.84	4.59	3.1	3.94	2.93		
NT2RP3000451	1.76	1.76	5.12	3.7	5	2.96		
NT2RP3000456	1.69	1.69	5.48	4.23	6.67	4.21		
NT2RP3000460	18.87	18.87	36.67	24.52	25.24	26.25		
NT2RP3000471	3.14	3.14	6.49	2.74	4.98	5.84		
NT2RP3000477	19.96	19.96	23.67	28.98	17.78	32.78		
NT2RP3000478	5.86	5.86	8.95	5.21	8.98	2.6		
NT2RP3000481	5.48	5.48	5.76	2.76	3.61	1.52	**	-
NT2RP3000484	3.51	3.51	4.26	2.32	2.55	1.76	*	-
NT2RP3000487	1.77	1.77	7.4	5.07	4.03	4.97		
NT2RP3000512	3.29	3.29	17.7	15.17	15.9	14.52		
NT2RP3000523	13.05	13.05	30.74	31.75	27.83	34.4		
NT2RP3000526	3.07	3.07	7.38	5.18	6.31	4.64		
NT2RP3000527	2.83	2.83	6.5	3.76	7.25	5.03		
NT2RP3000531	2.9	2.9	7.71	5.11	5.51	4.69		
NT2RP3000532	5.74	5.74	5.6	5.75	8.39	4.26		
NT2RP3000542	6.23	6.23	8.1	7.21	7.3	6.39		
NT2RP3000554	8.81	8.81	15.22	13.78	10.56	14.95		
NT2RP3000561	1.21	1.21	3.51	3.11	2.76	2.25		
NT2RP3000562	1.84	1.84	3.5	3.7	3.87	3.23		
NT2RP3000578	1.56	1.56	2.54	2.54	3.37	2.36		
NT2RP3000582	1.26	1.26	4.66	2.24	2.52	0.41		
NT2RP3000584	2.82	2.82	6.52	3.2	2.5	2.02		
NT2RP3000586	4.08	4.08	4.59	3.28	3.9	2.87		
NT2RP3000590	5.69	5.69	4.61	3.78	4.35	2.57		
NT2RP3000592	1.8	1.8	2.99	2.97	2.75	3.15		
NT2RP3000596	2.27	2.27	4.89	4.5	3.33	3.03		
NT2RP3000599	1.67	1.67	3.07	3.88	4.98	3.82	*	+
NT2RP3000603	6.09	6.09	39.25	40.43	44.88	35.89		
NT2RP3000605	2.84	2.84	6.66	4.56	4.23	2.56		
NT2RP3000607	5.35	5.35	7.59	5.74	8.46	7.55		
NT2RP3000616	3.26	3.26	5.45	2.56	2.38	1.21		
NT2RP3000621	5.18	5.18	8.48	10.28	10.29	6.01		
NT2RP3000622	2.36	2.36	8.76	5.85	6.21	4.72		
NT2RP3000624	1.53	1.53	3.19	3.97	3.06	2.78		
NT2RP3000628	2.44	2.44	8.04	10.27	7.85	5.58		
NT2RP3000631	4.71	4.71	14.95	22.82	16.45	14.2		
NT2RP3000632	2.35	2.35	5.5	7.78	8.91	5.91	*	+
NT2RP3000638	6.95	6.95	17.93	11.8	11.6	9.97		
NT2RP3000644	25.72	25.72	48.41	57.98	72.01	52.49	*	+
NT2RP3000645	5.85	5.85	10.48	9.84	12.55	8.43		
NT2RP3000652	3.39	3.39	5.34	6.22	5.9	7.74	*	+
NT2RP3000658	2.26	2.26	5.01	6.16	4.24	4.86		
NT2RP3000660	2.34	2.34	6.25	6.98	6.91	5.14		
NT2RP3000661	1.98	1.98	4.49	4.06	3.87	3.1		

(1130)

特開 2002-191363

2257

2258

NT2RP3000665	4.79	4.79	12.26	11.83	11.92	7		
NT2RP3000676	4.46	4.46	7.55	6.65	7.81	5.42		
NT2RP3000677	2.87	2.87	4.13	2.44	3.07	1.54		
NT2RP3000681	19.85	19.85	30.12	32.94	41.51	34.34	*	+
NT2RP3000683	2.68	2.68	9.67	6.69	7.09	6.69		
NT2RP3000685	1.7	1.7	2.5	3.63	2.36	3.44		
NT2RP3000690	2.77	2.77	3.29	3.82	3.75	2.72		
NT2RP3000698	10	10	22.49	25.66	17.08	27.43		
NT2RP3000708	3.45	3.45	5.5	8.17	9.22	8.56	**	+
NT2RP3000719	2.83	2.83	2.83	1.16	1.7	1.91	**	-
NT2RP3000721	5.63	5.63	24.61	23.43	39.76	21.55		
NT2RP3000728	3.33	3.33	2.57	1.4	1.64	1.05	**	-
NT2RP3000730	2.06	2.06	5.04	2.76	4.23	1.86		
NT2RP3000733	2.87	2.87	6.32	3.48	4.47	4.25		
NT2RP3000735	1.74	1.74	4.22	1.81	2.22	1.26		
NT2RP3000736	2.71	2.71	6.35	3.29	5.05	3.65		
NT2RP3000739	13.76	13.76	12.16	18.05	9.37	20.19		
NT2RP3000742	3.89	3.89	10.06	4.54	4.97	4.43		
NT2RP3000753	2.29	2.29	3.9	2.17	2.65	6.3		
NT2RP3000759	9.07	9.07	15.99	11.11	17.14	23.05		
NT2RP3000789	1.58	1.58	5.76	4.89	4.23	3.69		
NT2RP3000815	1.91	1.91	5.92	4.49	5.57	3.08		
NT2RP3000818	4.35	4.35	11.29	6.64	10.49	8.27		
NT2RP3000820	9.01	9.01	18.49	18.58	20.1	16.9		
NT2RP3000821	2.13	2.13	4.83	3.28	5.19	2.02		
NT2RP3000825	1.87	1.87	4.94	1.92	1.47	2.27		
NT2RP3000826	4.04	4.04	13.59	10.86	13.8	12.94		
NT2RP3000836	5.33	5.33	11.61	11.55	14.11	13.3		
NT2RP3000838	319.2	319.2	741.74	710.2	743.55	1049.86		
NT2RP3000839	2.35	2.35	6.67	4.53	6.38	4.36		
NT2RP3000841	2.17	2.17	4.32	3.79	5.55	4.72		
NT2RP3000845	3.96	3.96	8.89	5.76	6.71	7.85		
NT2RP3000847	3.7	3.7	7.94	4.48	5.94	5.28		
NT2RP3000848	2.84	2.84	8.34	5.36	6.81	6.3		
NT2RP3000850	5.67	5.67	7.04	6.58	11.29	7.47		
NT2RP3000852	3.27	3.27	3.17	4.02	5.23	5.8	*	+
NT2RP3000859	2.76	2.76	7.12	4.46	7.11	8.43		
NT2RP3000861	2.58	2.58	10.51	6.13	10.36	6.43		
NT2RP3000862	15.29	15.29	24.16	16.36	9.81	23.13		
NT2RP3000865	1.58	1.58	4.26	2.54	4.21	1.83		
NT2RP3000866	2.08	2.08	5.03	2.37	3.59	5.22		
NT2RP3000868	2.2	2.2	7.09	3.04	3.84	2.28		
NT2RP3000869	3.54	3.54	11.36	9.61	15.76	7.9		
NT2RP3000871	1.75	1.75	3.79	1.81	3.24	1.94		
NT2RP3000875	0.99	0.99	4.25	2.57	2.71	3.64		
NT2RP3000895	2.54	2.54	5.56	2.84	3.55	4.93		
NT2RP3000900	6.01	6.01	11.86	11.3	7.7	14.58		
NT2RP3000901	3.67	3.67	7.03	4.11	6.39	5.3		
NT2RP3000903	3.76	3.76	7.87	3.12	5.92	3.93		
NT2RP3000904	3.83	3.83	8.67	3.05	4.87	3.16		

(1131)

特開2002-191363

2259

2260

NT2RP3000907	5.66	5.66	10.03	8.94	10.67	10.14		
NT2RP3000913	6.04	6.04	15.01	17.87	25.57	17.37	*	+
NT2RP3000917	7.64	7.64	16.58	7.66	6.56	13.51		
NT2RP3000919	1.99	1.99	5.15	3.5	4.3	2.68		
NT2RP3000921	2.26	2.26	7.67	4.88	4.07	6.51		
NT2RP3000942	2.66	2.66	3.89	2.68	4.12	2.63		
NT2RP3000968	70.24	70.24	87.55	105.89	110.05	115.38	**	+
NT2RP3000974	5.36	5.36	9.06	3.21	4.48	2.64		
NT2RP3000980	5.77	5.77	5.77	2.09	4.14	2.26	*	-
NT2RP3000984	3.17	3.17	7.65	6.33	6.68	4.58		
NT2RP3000994	2.09	2.09	4.88	2.4	3.14	3.37		
NT2RP3001001	1.46	1.46	3.45	3.75	4.14	1.31		
NT2RP3001004	3.37	3.37	6.52	3.51	5.63	5.01		
NT2RP3001007	4.46	4.46	9.87	10.02	10.62	6.81		
NT2RP3001012	2.78	2.78	5.4	4.99	6.83	3.65		
NT2RP3001042	4.74	4.74	5.52	2.99	5.18	1.38		
NT2RP3001044	6.26	6.26	7.12	7.16	6.76	4.92		
NT2RP3001048	2.52	2.52	3.01	3.5	3.93	2.42		
NT2RP3001050	1.79	1.79	4.99	4.68	5.94	4.7		
NT2RP3001055	6.55	6.55	15.6	16.48	12.44	20.49		
NT2RP3001057	2.79	2.79	10.84	5.57	6.05	5.67		
NT2RP3001061	3.18	3.18	6.57	5.03	7.85	5.42		
NT2RP3001069	6.03	6.03	14.95	18.49	17.53	15.08		
NT2RP3001074	4.2	4.2	7.22	8.72	10	6.64		
NT2RP3001078	5.11	5.11	7.29	7.51	8.72	5.18		
NT2RP3001081	4	4	5.72	4.65	5.19	3.52		
NT2RP3001084	2.7	2.7	7.92	6.85	6.71	6.23		
NT2RP3001095	1.57	1.57	3.88	3.69	3.68	3.42		
NT2RP3001096	2.52	2.52	7.33	16.78	8.08	18.7	*	+
NT2RP3001097	3.65	3.65	4.28	6.42	8.11	8.5	**	+
NT2RP3001107	3.69	3.69	4.79	3.77	4.03	2.37		
NT2RP3001109	3.2	3.2	5.5	6.01	9.56	7.26	*	+
NT2RP3001111	4.58	4.58	4.19	3.41	3.51	2.29	*	-
NT2RP3001112	12.61	12.61	18.48	25.73	29.85	24.61	**	+
NT2RP3001113	1.21	1.21	2.59	2.47	3.24	2.19		
NT2RP3001115	1.51	1.51	3.32	2.57	3.77	2.19		
NT2RP3001116	1.01	1.01	2.66	2.55	4.4	2.91		
NT2RP3001119	3.69	3.69	6.75	9.07	6.67	5.44		
NT2RP3001120	5.02	5.02	8.24	8.85	7.87	6.71		
NT2RP3001126	6.16	6.16	12.34	17.84	19.66	17.49	**	+
NT2RP3001127	6.93	6.93	6.76	4.79	7.63	6.36		
NT2RP3001133	3.95	3.95	4.95	3.95	4.16	3.62		
NT2RP3001140	1.46	1.46	2.43	3.21	2.38	6.71		
NT2RP3001147	3.16	3.16	6.96	16.08	14.49	13.84	**	+
NT2RP3001150	1.99	1.99	4.32	4.06	5.68	3.76		
NT2RP3001152	1.7	1.7	3.29	3.15	3.62	2.51		
NT2RP3001155	2.95	2.95	4.35	3.68	4.35	3.58		
NT2RP3001156	4.38	4.38	6.57	2.91	5.72	5.67		
NT2RP3001159	5.38	5.38	10.5	7.87	10.86	7.25		
NT2RP3001170	7.38	7.38	5.96	6.12	8.01	4.56		

(1132)

特開 2002-191363

2261

2262

NT2RP3001176	3.49	3.49	10.75	6.27	8.23	9.49		
NT2RP3001195	2.35	2.35	4.81	6.79	5.79	6.22	*	+
NT2RP3001209	3.47	3.47	5.98	5.96	4.64	5.22		
NT2RP3001214	1.63	1.63	4.91	3.44	3.87	3.81		
NT2RP3001216	3.58	3.58	6.38	6.25	4.33	3.6		
NT2RP3001221	3.33	3.33	4.27	3.07	3.06	1.79		
NT2RP3001226	5.96	5.96	29.04	21.93	31.45	17.76		
NT2RP3001230	3.17	3.17	2.41	3.09	3.14	1.56		
NT2RP3001232	1.8	1.8	4.72	2.36	3.7	2.85		
NT2RP3001236	1.68	1.68	4.3	1.7	3.26	1.47		
NT2RP3001239	1.58	1.58	5.21	2.81	4.31	2.01		
NT2RP3001240	12.83	12.83	22.18	23.01	24.3	14.46		
NT2RP3001245	3.53	3.53	9.88	4.08	6.36	3.39		
NT2RP3001253	2.79	2.79	4.87	3.34	4.53	5.21		
NT2RP3001259	6.62	6.62	11.97	12.33	15.62	11.83		
NT2RP3001260	3.74	3.74	5.15	3.45	5.44	3.97		
NT2RP3001264	2.2	2.2	10.29	5.99	6.92	6.38		
NT2RP3001268	2.25	2.25	7.18	4.93	4.72	4.35		
NT2RP3001271	7.06	7.06	16.29	13.07	12.27	14.24		
NT2RP3001272	3.73	3.73	12.45	9.43	11.09	10.15		
NT2RP3001274	6.08	6.08	8.09	6.72	6.35	5.11		
NT2RP3001275	9.78	9.78	11.58	21.56	26.84	22.59	**	+
NT2RP3001280	3.39	3.39	5.5	3.58	5.24	4.18		
NT2RP3001281	3.15	3.15	3.89	3.08	4.48	5.14		
NT2RP3001288	49.31	49.31	103.24	124.07	142.92	164.41	*	+
NT2RP3001297	6.39	6.39	42.01	37.04	42.75	41.14		
NT2RP3001300	5.23	5.23	15.92	16.78	17.41	17.76		
NT2RP3001301	2.91	2.91	6.59	3.96	4.58	3.9		
NT2RP3001307	1.76	1.76	7.67	2.07	2.81	2.06		
NT2RP3001310	11.55	11.55	17.04	25.54	26.07	28.13	**	+
NT2RP3001318	2.11	2.11	3.4	2.49	3.37	2.37		
NT2RP3001322	3.58	3.58	5.23	2.62	3.84	5.48		
NT2RP3001325	2.7	2.7	8.39	5.82	6.82	5.58		
NT2RP3001338	2.67	2.67	6.19	4.1	4.21	3.5		
NT2RP3001339	2.53	2.53	5.64	3.08	4.89	2.91		
NT2RP3001340	2.9	2.9	8.42	6.36	7.07	5.79		
NT2RP3001341	2.26	2.26	6.97	5.1	5.62	4.73		
NT2RP3001354	3.22	3.22	9.77	4.28	6.93	9.35		
NT2RP3001355	1.9	1.9	5.41	2.65	3.82	2.74		
NT2RP3001356	2	2	5.34	2.59	3.2	3.55		
NT2RP3001359	1.09	1.09	4.05	1.63	2.5	1.75		
NT2RP3001364	2.34	2.34	5.31	3.26	6.67	2.67		
NT2RP3001373	1.12	1.12	3.22	2.1	3.74	1.71		
NT2RP3001374	1.9	1.9	4.17	3.18	3.92	3.1		
NT2RP3001383	3.84	3.84	8.96	3.92	6.65	3.85		
NT2RP3001384	4.11	4.11	9.47	3.54	4.46	2.41		
NT2RP3001388	3.98	3.98	8.79	9.48	10.99	9.4		
NT2RP3001392	4.61	4.61	6.19	3.91	6.14	3.23		
NT2RP3001396	1.7	1.7	6.39	4.04	4.66	4.53		
NT2RP3001398	2.51	2.51	6.55	3.85	7.05	2.94		

(1133)

特開2002-191363

2263

2264

NT2RP3001399	4.91	4.91	20.67	15.86	16.12	12.44		
NT2RP3001402	6.46	6.46	36.36	33.37	41.61	39.66		
NT2RP3001407	6.96	6.96	19.16	13.69	17.65	12.35		
NT2RP3001416	7.92	7.92	15.88	13.02	18.3	14.72		
NT2RP3001420	5.33	5.33	6.4	3.27	3.64	1.8	*	-
NT2RP3001425	3.73	3.73	4.92	4.74	5.67	3.15		
NT2RP3001426	2.39	2.39	6.08	5.45	4.45	5.11		
NT2RP3001427	1.82	1.82	5.61	3.46	2.89	3.59		
NT2RP3001428	2.42	2.42	6.29	5.69	4.81	3.77		
NT2RP3001429	3.08	3.08	5.91	4.15	7.37	4.73		
NT2RP3001432	2.14	2.14	6.61	3.72	4.44	3.58		
NT2RP3001439	4.14	4.14	6.39	5.87	7.27	4.41		
NT2RP3001441	6.45	6.45	12.63	11.13	14.61	11.2		
NT2RP3001446	4.99	4.99	4.99	4.64	5.22	4.39		
NT2RP3001447	2.72	2.72	5.21	6.64	5.14	6.33		
NT2RP3001449	3.95	3.95	11.85	16.9	14.57	13.16	*	+
NT2RP3001453	1.84	1.84	3.66	3.5	4.4	2.81		
NT2RP3001457	3.86	3.86	7.71	6.06	6.93	5.5		
NT2RP3001459	2.39	2.39	6.03	2.64	2.78	1.17		
NT2RP3001463	2.77	2.77	6.74	5.93	5.94	3.98		
NT2RP3001466	2.87	2.87	3.56	1.19	1.43	0.78	**	-
NT2RP3001472	5.74	5.74	4.02	3.7	4.85	4.32		
NT2RP3001475	3.54	3.54	7.61	6.91	6.65	7.39		
NT2RP3001479	2.54	2.54	6.66	4.37	5.69	5.16		
NT2RP3001490	3.18	3.18	9.26	4.4	6.02	5.21		
NT2RP3001492	4.36	4.36	7.84	7.59	7.08	5.72		
NT2RP3001495	4.14	4.14	3.85	2.75	2.92	1.76	*	-
NT2RP3001497	5.8	5.8	6.32	7.47	9.96	6.8		
NT2RP3001501	5.36	5.36	5.52	3.12	4.49	3.43	*	-
NT2RP3001527	4.89	4.89	6.71	4.9	5.14	3.52		
NT2RP3001529	1.51	1.51	3.5	4.12	3.95	4.18	*	+
NT2RP3001538	1.78	1.78	6.2	6.93	7.81	6.23		
NT2RP3001539	5.81	5.81	14.5	15.19	14.15	16.47		
NT2RP3001542	1.52	1.52	5.26	4.23	4.38	2.13		
NT2RP3001549	4.75	4.75	11.12	14.57	11.37	13.44		
NT2RP3001554	3.06	3.06	6.16	6.37	7.5	5.05		
NT2RP3001560	4.96	4.96	5.73	4.67	6.35	2.36		
NT2RP3001561	8.85	8.85	20.77	20.38	27.2	17.15		
NT2RP3001564	1.54	1.54	8.24	6.43	4.53	5.96		
NT2RP3001568	2.1	2.1	7.68	11.84	10.29	8.49	*	+
NT2RP3001575	3.94	3.94	7.24	6.39	6.97	6.16		
NT2RP3001580	1.78	1.78	4.49	4.35	3.8	3.11		
NT2RP3001587	4.38	4.38	8.74	10.75	10.04	7.77		
NT2RP3001589	3.17	3.17	8.21	5.6	7.79	4.36		
NT2RP3001592	4.52	4.52	21.6	19	32.62	14.54		
NT2RP3001607	3.42	3.42	1.86	1.59	2.8	1		
NT2RP3001608	1.05	1.05	3.59	2.41	1.73	2.31		
NT2RP3001613	3.08	3.08	2.77	3.89	2.91	3.99		
NT2RP3001619	4.31	4.31	8.15	7.69	6.45	7.62		
NT2RP3001621	1.18	1.18	2.69	2.39	2.28	2.02		

(1134)

特開2002-191363

2265

2266

NT2RP3001629	2.58	2.58	3.28	2.68	2.41	1.7		
NT2RP3001630	3.39	3.39	4.56	1.67	2.02	1.17	**	-
NT2RP3001631	9.01	9.01	14.34	18.65	21.16	15.24	*	+
NT2RP3001634	4	4	5.29	4.51	6.89	5.11		
NT2RP3001642	3.71	3.71	7.45	5.77	4.41	5.09		
NT2RP3001646	1.56	1.56	3.7	0.89	2.79	0.95		
NT2RP3001650	2.06	2.06	5.81	4.86	7.03	2.08		
NT2RP3001667	4.66	4.66	11.91	6.93	9.95	5.13		
NT2RP3001671	2.28	2.28	7.98	7.7	4.69	5.99		
NT2RP3001672	1.33	1.33	4.55	1.66	1.47	1.72		
NT2RP3001676	2.18	2.18	5.02	2.35	3.14	2.24		
NT2RP3001678	2.86	2.86	9.24	5.12	5.14	6.03		
NT2RP3001679	6.12	6.12	9.19	6.74	4.73	6.91		
NT2RP3001682	1.82	1.82	5.09	4.45	6.18	3.35		
NT2RP3001685	3.02	3.02	6.74	3.52	6.53	3.01		
NT2RP3001688	3.01	3.01	9.42	5.46	8.21	6.43		
NT2RP3001690	3.21	3.21	4.87	2.91	3.54	2.99		
NT2RP3001693	5.69	5.69	10.93	16.59	18.34	16.12	**	+
NT2RP3001696	2.28	2.28	3.63	1.77	3.68	3.39		
NT2RP3001698	35.35	35.35	79.65	85.09	91.88	105.32		
NT2RP3001708	4.82	4.82	8.78	6.34	6.95	9.01		
NT2RP3001712	8.69	8.69	16.06	10.22	14.19	13		
NT2RP3001716	1.44	1.44	5.45	2.14	3.42	2.31		
NT2RP3001724	2.75	2.75	6	4.08	4.54	2.63		
NT2RP3001727	11.73	11.73	38.73	39.17	49.36	31.26		
NT2RP3001729	3.36	3.36	4.7	5.69	6.55	3.06		
NT2RP3001730	12.54	12.54	26.52	12.53	19.94	16.4		
NT2RP3001733	1.46	1.46	3.04	2.09	3.7	1.62		
NT2RP3001737	3.02	3.02	7.12	4.62	5.49	2.78		
NT2RP3001738	1.59	1.59	8.22	3.38	6.01	3.03		
NT2RP3001739	3.26	3.26	5.25	5.63	6.1	2.51		
NT2RP3001742	2.54	2.54	5.36	3.86	4.55	4.03		
NT2RP3001751	3.61	3.61	11.54	9.94	12.82	8.76		
NT2RP3001752	2.58	2.58	7.01	2.1	3.59	2.76		
NT2RP3001753	5.73	5.73	9.48	10.83	15.3	13.69	*	+
NT2RP3001754	4.63	4.63	9.08	5.86	3.73	5.33		
NT2RP3001756	4.66	4.66	7.36	9.37	5.75	8.03		
NT2RP3001764	2.1	2.1	3.76	2.54	4.25	2.49		
NT2RP3001771	2.63	2.63	3.2	1.52	4.14	1.22		
NT2RP3001777	2.59	2.59	5.99	3.25	5.19	3.26		
NT2RP3001782	3.52	3.52	14.68	6.47	6.63	6.47		
NT2RP3001792	2.27	2.27	4.35	2.91	4.09	1.35		
NT2RP3001799	1.76	1.76	5.18	5.71	6.36	5.68		
NT2RP3001819	1.36	1.36	4.54	1.7	1.52	2.06		
NT2RP3001829	21.63	21.63	43.14	35.64	17.14	24.87		
NT2RP3001836	7.31	7.31	10.67	15.24	7.26	11.37		
NT2RP3001839	18.86	18.86	31.77	31.97	19.23	17.53		
NT2RP3001844	4.15	4.15	11.37	8.33	9.59	8.54		
NT2RP3001848	9.61	9.61	52.04	44.52	74.75	43.87		
NT2RP3001854	6.41	6.41	11.29	12.86	16.75	13.26	*	+

(1135)

特開 2002-191363

2267

2268

NT2RP3001855	2.27	2.27	3.94	1.87	1	1.74		
NT2RP3001857	3.1	3.1	5.22	5.13	5.6	3.33		
NT2RP3001858	1.53	1.53	4.45	4.41	6.04	2.97		
NT2RP3001861	7.35	7.35	16.34	11.85	14.46	10.42		
NT2RP3001866	4.35	4.35	9.63	5.52	10.42	7.93		
NT2RP3001871	4.82	4.82	6.34	5.55	6.38	4.55		
NT2RP3001874	6.8	6.8	9.73	7.72	11.19	6.14		
NT2RP3001878	5.98	5.98	6.35	4.59	6.89	5.02		
NT2RP3001885	3.21	3.21	5.3	5.65	5.25	4.42		
NT2RP3001896	1.64	1.64	3.49	3.37	2.13	2.02		
NT2RP3001898	9.03	9.03	17.69	14.71	8.69	11.94		
NT2RP3001899	3.21	3.21	7.33	4.52	6.9	5.81		
NT2RP3001901	4.58	4.58	9.18	8.19	9.44	9.21		
NT2RP3001915	4.84	4.84	11.12	14.09	15.67	14.04	*	+
NT2RP3001926	2.8	2.8	4.88	2.47	2.45	1.65		
NT2RP3001929	3.74	3.74	4.06	2.56	3.38	0.86		
NT2RP3001931	4.63	4.63	5.26	3.9	5.62	2.98		
NT2RP3001938	2.27	2.27	5.53	4.93	3.93	4.75		
NT2RP3001943	3.27	3.27	5.36	5.77	6.5	5.56		
NT2RP3001944	1.77	1.77	3.72	4.08	5.91	4.34		
NT2RP3001945	4.25	4.25	12.2	11.86	11.78	6.11		
NT2RP3001947	2.94	2.94	5.89	4.06	5.41	3.4		
NT2RP3001949	4.21	4.21	8.9	10.49	11.08	8.08		
NT2RP3001952	23.54	23.54	43.64	48.59	88.56	41.86		
NT2RP3001954	5.06	5.06	3.68	4.34	4.79	1.85		
NT2RP3001956	4.97	4.97	9.44	7.76	8.22	6.29		
NT2RP3001967	3.78	3.78	7.74	6.7	5.37	5.66		
NT2RP3001969	1.71	1.71	2.91	4.05	4.39	3.62	*	+
NT2RP3001976	2.25	2.25	4.67	6.22	6.25	4.84	*	+
NT2RP3001986	3.55	3.55	3.88	3.43	2.82	2.19		
NT2RP3001989	3.76	3.76	5.23	2.86	3.58	2.54		
NT2RP3002002	6.68	6.68	9.47	6.25	8.85	3.86		
NT2RP3002004	5.02	5.02	6.23	3.79	5.74	3.55		
NT2RP3002007	1.29	1.29	2.3	3.46	4.05	1.69		
NT2RP3002014	1.38	1.38	6.23	6.04	6.24	4.21		
NT2RP3002015	3.61	3.61	10.33	14.17	9.94	8.85		
NT2RP3002033	1.54	1.54	5.03	7.29	5.03	3.65		
NT2RP3002045	1.89	1.89	5.29	4.67	4.36	2.5		
NT2RP3002054	5.26	5.26	8.12	6.27	9.17	5.42		
NT2RP3002056	5.67	5.67	5.52	4.24	4.24	2.7	*	-
NT2RP3002057	4.35	4.35	3.5	2.87	2.41	0.81	*	-
NT2RP3002061	4.71	4.71	13.94	8.64	8.9	10.74		
NT2RP3002062	0.8	0.8	2.42	3.58	3.26	1.11		
NT2RP3002063	5.61	5.61	10.31	9.29	9.3	7.31		
NT2RP3002064	2.6	2.6	3.37	2.72	3.74	2.52		
NT2RP3002071	1.6	1.6	3.91	1.99	3.29	1.45		
NT2RP3002073	6.47	6.47	9.55	10.45	11.13	8.64		
NT2RP3002074	4.2	4.2	7.25	6.33	7.82	4.24		
NT2RP3002075	7.58	7.58	11.93	21.64	30.17	18.15	*	+
NT2RP3002077	3.81	3.81	5.95	2.48	3.05	2.78		

(1136)

特開 2002-191363

2269

2270

NT2RP3002081	4.25	4.25	7.55	13.22	12.62	11.13	**	+
NT2RP3002086	3.86	3.86	9.77	5.59	8.66	6.95		
NT2RP3002094	7.34	7.34	10.28	13.84	14.79	11.67	*	+
NT2RP3002096	1.98	1.98	4.53	1.28	3.12	1.73		
NT2RP3002097	3.77	3.77	6.16	6.1	8.34	6.88		
NT2RP3002098	1.61	1.61	4.3	1.04	1.8	1.46		
NT2RP3002102	2	2	4.86	3.11	3.4	3.16		
NT2RP3002106	2.74	2.74	4.98	2.83	4.9	2.51		
NT2RP3002108	3.69	3.69	7.8	3.11	3.39	3.15		
NT2RP3002109	12.49	12.49	32.04	31.61	27.15	25.12		
NT2RP3002110	36.38	36.38	54.93	55.24	58.94	46.55		
NT2RP3002113	11.15	11.15	13.99	10.66	15.22	11.44		
NT2RP3002120	2.22	2.22	4.42	2.31	4.13	2.7		
NT2RP3002121	5.93	5.93	14.39	13.38	14.39	15.06		
NT2RP3002126	34.03	34.03	108.96	121.18	130.55	142.49	*	+
NT2RP3002128	4.06	4.06	8.23	3.36	6.87	3.92		
NT2RP3002130	8.29	8.29	18.59	11.69	10.7	14.03		
NT2RP3002133	14.24	14.24	18.31	10.06	8.3	19.51		
NT2RP3002136	10.32	10.32	15.42	12.49	17.64	17.49		
NT2RP3002140	3.13	3.13	6.35	6.15	3.9	4.16		
NT2RP3002142	16.86	16.86	50.85	56.54	81.25	62.65	*	+
NT2RP3002146	4	4	7.22	5.14	9.31	6.56		
NT2RP3002147	3.8	3.8	10.45	6.06	7.4	6.2		
NT2RP3002151	5.62	5.62	10.64	6.27	4.43	7.33		
NT2RP3002155	1.62	1.62	3.27	2.01	4.98	0.8		
NT2RP3002156	3.15	3.15	4.82	2.55	4.08	2.42		
NT2RP3002160	1.57	1.57	3.43	1.36	3.43	1.56		
NT2RP3002163	20.86	20.86	55.1	35.13	44.03	32.6		
NT2RP3002165	4.17	4.17	3.67	6.21	8.31	4.86		
NT2RP3002166	4.04	4.04	10.53	7.76	8.79	5.58		
NT2RP3002173	2.24	2.24	5.75	2.95	3.34	3.53		
NT2RP3002174	8.41	8.41	15.8	13.21	7.82	14.77		
NT2RP3002181	1.1	1.1	3.46	1.87	3.51	1.61		
NT2RP3002185	2.69	2.69	4.51	2.94	4.35	2.61		
NT2RP3002193	5.51	5.51	13.38	16.39	15.35	11.36		
NT2RP3002204	5.66	5.66	12.49	17.04	24.14	18.95	*	+
NT2RP3002244	4.03	4.03	8.29	5.28	6.11	4.8		
NT2RP3002248	5.42	5.42	11.1	8.19	11.78	6.52		
NT2RP3002253	2.61	2.61	9.3	9.66	11.26	6.18		
NT2RP3002255	11.07	11.07	26.56	22.78	11.53	20.93		
NT2RP3002264	3.06	3.06	5.54	5.88	7.37	4.07		
NT2RP3002267	1.26	1.26	4.33	3.1	4.65	1.82		
NT2RP3002273	7.51	7.51	12.98	10.15	13.8	12.11		
NT2RP3002276	5.22	5.22	7.89	3.08	7.68	3.48		
NT2RP3002281	6.37	6.37	6.83	7.45	8.46	3.44		
NT2RP3002286	3	3	4.79	3.54	4.34	3.88		
NT2RP3002297	10.62	10.62	29.36	22.26	20.57	23.93		
NT2RP3002301	5.73	5.73	13.24	9.47	7.55	6.21		
NT2RP3002303	3.01	3.01	6.39	5.29	6.65	4.58		
NT2RP3002304	2.66	2.66	7.17	6.3	7.3	4.91		

(1137)

特開 2002-191363

2271

2272

NT2RP3002309	2.3	2.3	7.18	9.26	13	4.39		
NT2RP3002311	4.54	4.54	6.67	3.17	4.02	1.83		
NT2RP3002315	15.27	15.27	20.91	25.82	33.13	21.82		
NT2RP3002319	2.37	2.37	5.06	3.07	3.51	2.38		
NT2RP3002324	8.97	8.97	61.42	49.85	51.23	55.4		
NT2RP3002330	4.74	4.74	8.33	10.31	8.24	8.15		
NT2RP3002333	5.13	5.13	14.32	13.14	13.65	8.12		
NT2RP3002337	2.61	2.61	5.14	4.8	5.57	4.87		
NT2RP3002342	5.16	5.16	11.56	5.52	7.51	6.43		
NT2RP3002343	3.38	3.38	7.29	5.8	7.03	4.13		
NT2RP3002351	4.32	4.32	4.55	4.38	3.9	2.55		
NT2RP3002352	6.3	6.3	8.01	4.4	6.76	4.31		
NT2RP3002353	3	3	4.85	4.87	6.18	5.9		
NT2RP3002362	5	5	11.74	15.86	11.03	10.49		
NT2RP3002363	2.41	2.41	3.67	5.53	6.17	2.32		
NT2RP3002377	2.61	2.61	5.47	6.8	7.31	4.73		
NT2RP3002377	4.47	4.47	7.73	11.4	5.31	7.09		
NT2RP3002394	5.58	5.58	7.35	7.82	10.17	4.46		
NT2RP3002397	3.77	3.77	4.81	2.7	3.12	1.68	*	-
NT2RP3002399	4.61	4.61	7.69	14.65	13.02	16.16	**	+
NT2RP3002402	2.84	2.84	6.99	8.94	8.7	6.99		
NT2RP3002404	2.88	2.88	5.6	3.12	3.73	1.83		
NT2RP3002410	4.85	4.85	15.65	17.05	14.13	10.65		
NT2RP3002411	2.98	2.98	5.68	3.7	5.29	2.85		
NT2RP3002414	5.62	5.62	9.35	10.28	6.81	7.92		
NT2RP3002430	5.11	5.11	14.63	18.24	19.29	14.51		
NT2RP3002448	5.4	5.4	4.6	4.35	5.25	3.62		
NT2RP3002454	7.3	7.3	15.31	12.9	12.71	9.32		
NT2RP3002455	4.62	4.62	12.11	14.31	9.11	13.25		
NT2RP3002456	3.21	3.21	7.75	7.09	6.57	5.9		
NT2RP3002462	2.79	2.79	4.16	4.94	6.17	4.79	*	+
NT2RP3002469	3.84	3.84	6.38	9.24	6.78	8.07	*	+
NT2RP3002470	6.7	6.7	14.71	17.86	16.99	13.48		
NT2RP3002484	4.01	4.01	6.86	6.81	8.01	4.59		
NT2RP3002491	3.62	3.62	4.1	2.05	2.51	1.46	**	-
NT2RP3002494	79.24	79.24	131.02	118.47	163.2	105.2		
NT2RP3002497	1.07	1.07	1.57	2.37	1.39	1.37		
NT2RP3002500	1.23	1.23	1.13	2.72	2.02	2.07	**	+
NT2RP3002501	5.25	5.25	8.49	8.45	8.11	9.69		
NT2RP3002512	2.85	2.85	3.97	3.3	2.74	3.57		
NT2RP3002529	3.94	3.94	7.5	6.59	5.14	5.85		
NT2RP3002533	7.95	7.95	10.26	9.79	10.51	8.18		
NT2RP3002539	4.39	4.39	4.32	5.66	6.61	2.85		
NT2RP3002540	5.24	5.24	5.5	3.48	4.76	3.65	*	-
NT2RP3002543	3.44	3.44	7.17	4.93	6.21	5.31		
NT2RP3002545	7.34	7.34	7.46	5.17	5.52	6.8	*	-
NT2RP3002549	3.27	3.27	7.8	4.98	6.11	4.54		
NT2RP3002552	3.05	3.05	6.04	4.17	5.81	4.06		
NT2RP3002558	9.54	9.54	9.39	9.93	4.26	11.27		
NT2RP3002565	1.94	1.94	4.83	1.73	2.48	1.52		

(1138)

特開 2002-191363

2273

2274

NT2RP3002566	3.62	3.62	7.02	4.03	8.51	3.65		
NT2RP3002571	2.53	2.53	4.85	3.77	5.41	3.74		
NT2RP3002572	2.98	2.98	5.28	4.75	4.74	5.21		
NT2RP3002573	4.31	4.31	11.38	7.06	9.48	7.06		
NT2RP3002577	1.57	1.57	4.61	2.71	2.32	1.9		
NT2RP3002579	3.92	3.92	6.41	4.03	7.75	11.16		
NT2RP3002582	5.02	5.02	7.17	11.51	14.07	8.45	*	+
NT2RP3002587	1.9	1.9	3.13	2.68	3.04	1.97		
NT2RP3002590	3.16	3.16	5.65	9.06	10.39	8.06	**	+
NT2RP3002602	3.02	3.02	4.24	3.95	5.85	3.77		
NT2RP3002603	71.53	71.53	214.41	268.41	257.84	298.26	*	+
NT2RP3002621	1.95	1.95	3.42	2.13	5.13	1.85		
NT2RP3002622	2.63	2.63	7.38	3.7	7.36	4.67		
NT2RP3002624	2.29	2.29	7.4	4.04	4.9	3.64		
NT2RP3002628	6.36	6.36	16.17	19.57	22.15	16.21		
NT2RP3002629	8.96	8.96	13.58	15.4	18.26	15.57	*	+
NT2RP3002631	1.95	1.95	1.67	0.91	1.65	1.69		
NT2RP3002647	4.04	4.04	4.01	5.44	7.16	4.6		
NT2RP3002649	2.99	2.99	5.99	2.15	5.84	3.23		
NT2RP3002650	3.32	3.32	11.62	4.98	8.53	6.88		
NT2RP3002652	2.27	2.27	6.59	4.91	6.83	4.25		
NT2RP3002654	3.05	3.05	7.5	5.58	5.4	4.2		
NT2RP3002657	14.14	14.14	13.87	17.27	26.08	18.87		
NT2RP3002659	1.92	1.92	6.01	3.91	5.78	3.47		
NT2RP3002660	3.09	3.09	4.84	3.77	7.72	3.52		
NT2RP3002663	2.39	2.39	3.33	2.54	3.13	2.84		
NT2RP3002664	2.74	2.74	7.28	3.65	2.56	2.13		
NT2RP3002667	2.92	2.92	6.59	6.63	5.46	5.35		
NT2RP3002671	2.37	2.37	5.02	3.91	5.52	4.11		
NT2RP3002682	6.34	6.34	20.62	14.37	17.64	21.7		
NT2RP3002684	4	4	6.34	3.32	6.16	3.18		
NT2RP3002687	3.25	3.25	6.22	2.7	3.87	6.41		
NT2RP3002688	3.22	3.22	4.98	2.63	3.91	2.61		
NT2RP3002698	2.2	2.2	3.99	3.07	4.28	2.38		
NT2RP3002701	2.93	2.93	6.73	3.45	3.07	3.6		
NT2RP3002705	2.17	2.17	8.01	4.36	8.76	4.72		
NT2RP3002708	3.69	3.69	9.88	5.64	7.34	4.9		
NT2RP3002711	6.67	6.67	7.85	7.77	7.56	6.69		
NT2RP3002712	55.99	55.99	75.28	146.74	168.42	130.64	**	+
NT2RP3002713	4.31	4.31	7.06	2.66	2.19	1.87	*	-
NT2RP3002721	5.77	5.77	10.06	11.06	16.94	8.96		
NT2RP3002722	7.11	7.11	10.08	7.8	6.45	6.62		
NT2RP3002723	42.31	42.31	75.85	60.39	46.74	58.76		
NT2RP3002737	8.35	8.35	18.1	10.97	11.6	9.37		
NT2RP3002738	1.9	1.9	6.13	3.09	5.23	3.54		
NT2RP3002742	14.11	14.11	23.22	30.39	28.27	27.66	*	+
NT2RP3002744	4.09	4.09	5.24	3.92	4.92	1.71		
NT2RP3002756	5.8	5.8	5.8	3.19	2.68	1.55	**	-
NT2RP3002757	12	12	17.79	19.76	24.24	19.75	*	+
NT2RP3002758	21.11	21.11	42.35	44.47	63.91	36.38		

(1139)

特開2002-191363

2275

2276

NT2RP3002762	5.07	5.07	8.82	7.21	7.43	7.76		
NT2RP3002763	1.62	1.62	4.86	3.76	4.99	2.18		
NT2RP3002770	1.78	1.78	5.14	3.46	3.7	2.93		
NT2RP3002771	17.04	17.04	39.53	24.93	40.21	34.4		
NT2RP3002785	2.42	2.42	5.45	3.36	4.09	2.66		
NT2RP3002790	4.65	4.65	4.22	3.16	3.57	2.33	*	-
NT2RP3002799	4.73	4.73	6.33	3.42	2.7	1.43	*	-
NT2RP3002801	4.14	4.14	3.59	3.6	3.22	2.49		
NT2RP3002802	2.31	2.31	6.3	6.78	5.43	4.4		
NT2RP3002810	2.98	2.98	5.41	7.44	12.32	13.27	*	+
NT2RP3002818	1.5	1.5	2.44	2.18	4.16	2.47		
NT2RP3002821	12.8	12.8	33.14	26.1	35.81	23.02		
NT2RP3002823	3.85	3.85	8.98	4.65	5.92	3.87		
NT2RP3002825	5.47	5.47	13.04	13.47	19.19	7.12		
NT2RP3002829	5.37	5.37	6.25	4.75	4.89	3.8		
NT2RP3002831	4.01	4.01	6.13	9.07	8.77	5.19		
NT2RP3002836	7.33	7.33	19.42	11.56	16.91	20.66		
NT2RP3002845	4.17	4.17	6.63	7.87	8.6	8.45	*	+
NT2RP3002852	3.37	3.37	7.57	7.8	8.72	8.21		
NT2RP3002861	3.82	3.82	6.4	7.34	7.35	4.63		
NT2RP3002869	3.66	3.66	3.26	2.49	1.86	0.49	*	-
NT2RP3002874	11.25	11.25	21.44	25.33	31.95	25.54	*	+
NT2RP3002876	6.98	6.98	11.06	12.8	14.93	14.39	*	+
NT2RP3002877	4.7	4.7	5.96	3.3	5.24	2.53		
NT2RP3002887	0.47	0.47	3.42	2.81	3.53	3.91		
NT2RP3002900	6.46	6.46	19.64	21.86	21.54	22.3		
NT2RP3002902	4.01	4.01	10.25	11.72	8.52	8.06		
NT2RP3002909	2.61	2.61	6.19	6.67	5.38	3.93		
NT2RP3002911	3.05	3.05	3.68	3.09	3.73	2.24		
NT2RP3002948	4.09	4.09	4.81	2.73	4.44	2.07		
NT2RP3002953	3.85	3.85	3.6	2.27	2.52	0.84	*	-
NT2RP3002955	6.55	6.55	3.78	1.93	2.47	0.86	*	-
NT2RP3002958	5.85	5.85	11.4	16.5	10.57	16.64		
NT2RP3002969	4.28	4.28	8.27	12.91	7.49	7.33		
NT2RP3002972	3.55	3.55	4.82	4.41	6.18	2.29		
NT2RP3002978	1.48	1.48	2.99	2.61	3.7	1.49		
NT2RP3002983	2.89	2.89	4.69	4.46	6.12	4.37		
NT2RP3002985	4.23	4.23	17.87	13.64	20.26	11.12		
NT2RP3002988	3.97	3.97	4.6	4.12	5.64	4.13		
NT2RP3003000	3.11	3.11	3.46	2.46	3.2	1.51		
NT2RP3003008	3.26	3.26	5.87	3.95	4.55	2.96		
NT2RP3003012	3.43	3.43	6.06	3.9	4.96	2.79		
NT2RP3003015	1.35	1.35	4.9	1.5	2.5	0.54		
NT2RP3003018	2.15	2.15	6.09	3.45	7.24	2.59		
NT2RP3003028	3.53	3.53	7.23	3.5	5.05	4.01		
NT2RP3003029	111.75	111.75	149.73	175.13	159.77	181.4	*	+
NT2RP3003032	7.06	7.06	9.05	11.87	18.84	9.94		
NT2RP3003041	2.07	2.07	1.88	1.61	1.41	0.69		
NT2RP3003044	3.06	3.06	7.45	5.72	6.11	7.57		
NT2RP3003047	3.09	3.09	5.16	2.4	4.67	2.06		

(1140)

特開2002-191363

2277

2278

NT2RP3003050	5.96	5.96	12.03	6.74	10.3	8.42		
NT2RP3003053	7.46	7.46	18	14.42	17.14	13.72		
NT2RP3003059	1.93	1.93	4.76	2.88	4.41	3.06		
NT2RP3003061	2.8	2.8	8.59	5.49	5.68	5.16		
NT2RP3003068	5.99	5.99	11.77	9.41	8.75	10.04		
NT2RP3003071	7.22	7.22	10.77	10.39	14.52	10.39		
NT2RP3003076	2.67	2.67	9.49	6.57	6.57	4.01		
NT2RP3003078	1.5	1.5	4.12	2.09	4.52	2.43		
NT2RP3003081	6.21	6.21	10.54	9.94	9.6	9.97		
NT2RP3003090	1.49	1.49	5.95	3.28	3.57	3.25		
NT2RP3003097	2.42	2.42	7.15	2.71	3.72	2.93		
NT2RP3003098	2.75	2.75	4.22	2.73	3.43	1.73		
NT2RP3003101	5.56	5.56	7.24	7.73	10.35	7.18		
NT2RP3003109	16.11	16.11	27.38	27.36	41.03	21.91		
NT2RP3003121	3.39	3.39	11.03	4.61	8	2.44		
NT2RP3003133	2.09	2.09	5.78	4.93	8.58	3.96		
NT2RP3003137	3.42	3.42	5.74	6.32	7.59	5.29		
NT2RP3003138	2.36	2.36	5.27	5.26	5.7	3.74		
NT2RP3003139	2.53	2.53	7.8	3.15	5.74	2.7		
NT2RP3003145	5.08	5.08	32.56	25.8	29.74	19.72		
NT2RP3003150	2.03	2.03	5.17	3.56	3.76	1.97		
NT2RP3003157	2.52	2.52	8.34	6.4	10.1	4.94		
NT2RP3003185	1.77	1.77	3.88	1.91	3.34	2.62		
NT2RP3003193	2.62	2.62	5.75	6.03	4.59	2.65		
NT2RP3003197	2.38	2.38	3.8	3.11	4.02	2.2		
NT2RP3003203	11.82	11.82	14.35	16.85	10.17	15.27		
NT2RP3003204	3.76	3.76	7.93	4.04	6.17	3.79		
NT2RP3003210	14.48	14.48	75.3	58.97	84.6	68.66		
NT2RP3003212	5.15	5.15	9.44	9.21	10.67	7.36		
NT2RP3003213	4.16	4.16	5.68	5.15	7.02	5.44		
NT2RP3003224	1.7	1.7	4.75	2.43	2.11	2.64		
NT2RP3003226	3.25	3.25	5.68	6.57	5.94	3.63		
NT2RP3003230	7.79	7.79	11.47	12.39	8.89	6.72		
NT2RP3003235	7.61	7.61	10.79	7.77	7.73	6.89		
NT2RP3003242	12.17	12.17	23.49	26.68	32.03	19.25		
NT2RP3003251	5.61	5.61	9.47	3.73	4.95	4.08		
NT2RP3003252	3.95	3.95	5.95	2.19	3.7	2.42		
NT2RP3003258	4.92	4.92	7.89	19.94	24.95	15.47	**	+
NT2RP3003260	4.54	4.54	12.34	13.46	11.52	12.68		
NT2RP3003264	1.64	1.64	5.99	3.18	4.32	1.86		
NT2RP3003273	2.18	2.18	4.93	4.57	3.58	1.72		
NT2RP3003278	1.33	1.33	4	1.31	5.12	0.63		
NT2RP3003280	9.85	9.85	23.11	18.18	19.52	18.19		
NT2RP3003282	5.29	5.29	6.25	3.62	3.97	3.48	**	-
NT2RP3003290	6.64	6.64	9.09	4.8	5.38	3.78	*	-
NT2RP3003301	4.01	4.01	5.73	4.31	4.59	3.23		
NT2RP3003302	1.45	1.45	2.31	2.91	2.64	1.91		
NT2RP3003311	2.45	2.45	6.76	15.72	13.09	11.55	**	+
NT2RP3003312	1.81	1.81	3.35	3.73	3.87	2.41		
NT2RP3003313	1.61	1.61	4.2	2.91	5.4	2.87		

(1141)

特開2002-191363

2279

2280

NT2RP3003327	1.62	1.62	6.24	4.81	4.95	3.34		
NT2RP3003330	5.13	5.13	8.01	15.68	16.13	12.78	**	+
NT2RP3003344	3.36	3.36	4.14	2.92	3.74	2.6		
NT2RP3003346	3.81	3.81	4.83	4.38	4.05	1.24		
NT2RP3003349	4.04	4.04	6.93	9.96	9.41	9.65	**	+
NT2RP3003353	1.95	1.95	3.24	4.06	5.37	2.45		
NT2RP3003354	5.09	5.09	13.72	16.29	12.02	13.5		
NT2RP3003368	3.03	3.03	4.73	4.04	4.08	2.63		
NT2RP3003375	4.1	4.1	7.4	7.41	9.67	6.62		
NT2RP3003377	4.16	4.16	3.98	2.57	3.58	1.65		
NT2RP3003384	5.77	5.77	4.55	2.83	3.43	2.56	**	-
NT2RP3003385	4.55	4.55	3.12	1.9	2.36	1.47	*	-
NT2RP3003396	3.93	3.93	13.63	16.4	8.38	12.32		
NT2RP3003403	1.62	1.62	2.54	3.24	4.73	1.92		
NT2RP3003409	1.18	1.18	2.97	3.3	4.48	3.03		
NT2RP3003411	4.59	4.59	15.42	14.11	15.42	10.96		
NT2RP3003420	3.79	3.79	4.36	3.68	2.13	1.85		
NT2RP3003425	3.25	3.25	6.71	5.85	7.25	5.49		
NT2RP3003426	9.11	9.11	16.3	10.88	11.12	17.45		
NT2RP3003427	5.95	5.95	10.09	9.15	13.58	8.03		
NT2RP3003433	2.55	2.55	6.26	8.42	9.57	4.87		
NT2RP3003437	22.12	22.12	49.85	51.81	44	38.77		
NT2RP3003448	1.88	1.88	4.24	3.5	3.83	2.63		
NT2RP3003455	5.23	5.23	12.16	11.8	9.96	8.44		
NT2RP3003462	4.96	4.96	10.07	10.76	8.25	7.08		
NT2RP3003464	3.79	3.79	5.03	3.01	4.76	1.2		
NT2RP3003469	4.1	4.1	7.77	6.62	7.56	5.07		
NT2RP3003473	22.06	22.06	36.6	54.82	69.25	56.46	**	+
NT2RP3003474	8.26	8.26	23.04	13.23	12.04	13.52		
NT2RP3003475	2.84	2.84	4.04	4.55	4.45	3.28		
NT2RP3003490	2.7	2.7	5.81	4.21	4.43	3.12		
NT2RP3003491	2.26	2.26	3	2.14	2.75	1.26		
NT2RP3003493	11.75	11.75	30.77	34.59	28.9	34.45		
NT2RP3003500	4.93	4.93	5.26	4.99	7.46	3.65		
NT2RP3003527	2.73	2.73	3.09	2.42	2.92	1.72		
NT2RP3003532	2.7	2.7	1.81	2.14	3.33	2.13		
NT2RP3003535	3.14	3.14	4.37	1.92	3.4	2.19		
NT2RP3003536	3.04	3.04	5.95	3.45	5.2	4.97		
NT2RP3003543	2.61	2.61	6.24	3.21	4.81	3.83		
NT2RP3003549	1.43	1.43	6.66	2.18	4.07	1.55		
NT2RP3003552	1.8	1.8	5.76	0.64	0.84	1.58		
NT2RP3003555	4.4	4.4	14.14	12.16	17.43	16.23		
NT2RP3003559	2.81	2.81	6.7	3.88	5.11	5.49		
NT2RP3003564	3.11	3.11	5.9	2.24	4.6	5.25		
NT2RP3003572	2.1	2.1	4.21	1.88	3.02	2.32		
NT2RP3003576	5.88	5.88	10.15	11.32	8.98	9.22		
NT2RP3003587	7.39	7.39	12.41	10.01	12.71	12.75		
NT2RP3003589	15.33	15.33	22.45	23.89	23.75	26.58		
NT2RP3003592	7.77	7.77	10.4	8.42	14.48	9.74		
NT2RP3003593	8.16	8.16	13.62	13.47	13.84	110.49		

(1142)

特開2002-191363

2281

2282

NT2RP3003614	2.66	2.66	8.18	3.11	4.48	7.09		
NT2RP3003621	1.64	1.64	3.91	2.1	3.68	2.96		
NT2RP3003625	1.54	1.54	6.94	3.79	5.09	4.96		
NT2RP3003627	6.73	6.73	20.05	16.23	13.97	25.71		
NT2RP3003636	3.3	3.3	7.74	5.99	3.79	10.4		
NT2RP3003642	7.12	7.12	12.2	12.85	13.15	15.83		
NT2RP3003645	2.91	2.91	6.07	2.23	2.42	3.53		
NT2RP3003648	2.88	2.88	3.71	2.17	2.44	3.13		
NT2RP3003649	2.7	2.7	9.28	6.36	5.11	12.04		
NT2RP3003650	2.65	2.65	4.25	4.38	3.16	4.09		
NT2RP3003656	1.69	1.69	3.23	1.94	4.12	3		
NT2RP3003659	2.76	2.76	4.56	2.14	4.8	4.88		
NT2RP3003662	31.39	31.39	53.28	34.35	14.68	34.64		
NT2RP3003664	3.56	3.56	6.5	6.18	5.45	6.55		
NT2RP3003665	1.89	1.89	4.83	2.07	2.8	4.96		
NT2RP3003671	2.88	2.88	4.33	3.03	2.6	4.29		
NT2RP3003672	4.78	4.78	9.8	10.69	14.73	16.35	*	+
NT2RP3003673	4.98	4.98	9.42	5.35	3.05	4.12		
NT2RP3003679	40.1	40.1	95.75	69.92	23.86	83.88		
NT2RP3003680	3.13	3.13	5.38	3.96	4.58	5.88		
NT2RP3003686	2.22	2.22	4.43	2.84	4.85	2.25		
NT2RP3003689	4.05	4.05	9.69	5.94	5.63	8.27		
NT2RP3003697	13.79	13.79	120.74	108.93	77.49	68.74		
NT2RP3003701	2.7	2.7	5.17	2.58	3.05	2.57		
NT2RP3003704	2.99	2.99	6.96	7.09	7.61	6.96		
NT2RP3003714	1.39	1.39	4.25	1.68	0.89	1.14		
NT2RP3003716	2.05	2.05	4.23	3	2.29	2.24		
NT2RP3003721	1.83	1.83	3.27	1.85	3.45	2.18		
NT2RP3003722	3.45	3.45	8.18	8.08	7.79	5.45		
NT2RP3003726	3.5	3.5	4.9	2.77	4.51	2.32		
NT2RP3003729	4.1	4.1	8.53	4.22	5.44	4.6		
NT2RP3003731	5.06	5.06	6.98	4.19	3.54	7.45		
NT2RP3003740	2.58	2.58	5.08	2.42	2.48	2.94		
NT2RP3003746	3.63	3.63	8.14	6.7	5.94	6.59		
NT2RP3003749	0.67	0.67	2.58	1.55	2.08	1.73		
NT2RP3003754	3.32	3.32	7.31	4.66	5.87	5.81		
NT2RP3003759	1.16	1.16	4.44	2.49	4.41	1.43		
NT2RP3003764	3.97	3.97	7.08	6.85	7.41	5.06		
NT2RP3003766	6.93	6.93	7.84	3.3	5.87	3.79	*	-
NT2RP3003767	11.19	11.19	16.8	14.83	21.08	16.97		
NT2RP3003778	3.36	3.36	4.89	4.46	5.55	4.27		
NT2RP3003779	4.05	4.05	15.26	13.02	8.74	10.52		
NT2RP3003783	9.25	9.25	21.72	22.42	13.65	18.76		
NT2RP3003787	2.15	2.15	4.65	4.41	4.74	6.37		
NT2RP3003789	5.12	5.12	10.16	11.63	12.19	14.96	*	+
NT2RP3003795	1.48	1.48	6.48	4.09	2.82	2.24		
NT2RP3003799	2.67	2.67	5.5	3.08	2.38	1.75		
NT2RP3003800	4.36	4.36	5.92	4.14	4.57	6.91		
NT2RP3003805	8.15	8.15	6.78	8.4	5.48	5.89		
NT2RP3003809	1.94	1.94	7.2	5.83	5.4	4.82		

(1143)

特開 2002-191363

2283

2284

NT2RP3003819	3.39	3.39	6.07	7.3	5.97	6.35		
NT2RP3003824	5.69	5.69	10.69	14.08	14.85	13.32	*	+
NT2RP3003825	9.06	9.06	16.31	12.87	16.88	16.75		
NT2RP3003828	4.7	4.7	14.38	13.36	15.69	14.55		
NT2RP3003831	4.01	4.01	6.38	5.77	6.54	7.23		
NT2RP3003833	5.12	5.12	7.5	6.44	8.88	6.96		
NT2RP3003836	6.37	6.37	5.05	5.74	6.47	4.31		
NT2RP3003842	2.7	2.7	9.08	6.84	6.51	7.09		
NT2RP3003843	9.26	9.26	26.77	16.67	12.71	16.2		
NT2RP3003844	20.38	20.38	46.56	42.84	27.94	44.32		
NT2RP3003846	4.04	4.04	8.45	8.94	7.18	8.05		
NT2RP3003849	2.27	2.27	2.68	2.67	2.73	1.68		
NT2RP3003862	28.91	28.91	45.63	32	37.58	44.88		
NT2RP3003870	4.76	4.76	4.81	2.54	2.93	2.05	**	-
NT2RP3003874	21.46	21.46	20.88	33.11	47.25	36.44	*	+
NT2RP3003876	1.62	1.62	8.08	5.45	7.49	6.81		
NT2RP3003880	1.74	1.74	4.63	5.31	4.66	4.73		
NT2RP3003889	1.69	1.69	3.04	3.41	3.53	9.53		
NT2RP3003891	1.88	1.88	2.98	2.56	3.19	1.37		
NT2RP3003914	3.1	3.1	7.35	6.88	5.15	7.39		
NT2RP3003915	5.03	5.03	8.44	9.52	11.35	8.6		
NT2RP3003918	6.79	6.79	10.39	10.04	13.71	12.42		
NT2RP3003920	6.9	6.9	9.13	8.31	10.22	8.96		
NT2RP3003924	2.25	2.25	9.57	6.49	5.34	6.91		
NT2RP3003932	1.41	1.41	3.85	5.17	5.26	3.85		
NT2RP3003939	3.48	3.48	11.88	9.86	14.05	11.09		
NT2RP3003940	11.34	11.34	27.33	23.54	20.59	23.06		
NT2RP3003943	2.6	2.6	2.83	2.85	2.78	3.88		
NT2RP3003959	3.52	3.52	6.96	6.54	5.93	5.49		
NT2RP3003963	4.83	4.83	7.59	4.01	4.61	2.52		
NT2RP3003965	11.14	11.14	13.85	18.75	20.67	17.22	**	+
NT2RP3003972	26.1	26.1	40.32	22.13	17.98	38.48		
NT2RP3003973	2.85	2.85	4.33	1.96	3.6	3.96		
NT2RP3003979	5.89	5.89	12.53	6.92	8.49	8.84		
NT2RP3003980	3.52	3.52	9.41	9.34	8.89	7.92		
NT2RP3003982	4.2	4.2	4.63	2.44	1.6	4.61		
NT2RP3003989	6.24	6.24	4.69	9.61	5.62	16.05		
NT2RP3003992	2.13	2.13	4.89	2.47	5.12	4.8		
NT2RP3004000	2.81	2.81	6	1.72	3.22	2.62		
NT2RP3004001	11.38	11.38	19.94	11.62	11.37	21.11		
NT2RP3004005	2.89	2.89	7.79	4.7	4.48	6.84		
NT2RP3004013	2.23	2.23	7.2	2.66	4.87	3.57		
NT2RP3004016	1.5	1.5	7.1	2.22	3.14	2.88		
NT2RP3004025	4.02	4.02	7.69	7.48	12.19	9.01		
NT2RP3004030	7.05	7.05	12.64	13.97	15.8	17.66	*	+
NT2RP3004041	5.65	5.65	11.38	10.48	9.57	19.81		
NT2RP3004042	15.22	15.22	102.33	97.27	103.6	99.67		
NT2RP3004044	2.13	2.13	6.51	5.14	7.21	4.22		
NT2RP3004051	2.6	2.6	5.79	2.23	5.51	4.69		
NT2RP3004052	7.1	7.1	11.22	5.63	4.98	9.78		

(1144)

特開 2002-191363

2285

2286

NT2RP3004053	15.87	15.87	35.04	23.12	40.67	40.17		
NT2RP3004055	2.38	2.38	5.33	2.98	3.3	4.47		
NT2RP3004059	4.05	4.05	8.8	8.15	7.03	11		
NT2RP3004063	5.13	5.13	11.23	8.78	11.27	12.33		
NT2RP3004067	4.24	4.24	8.4	6.62	6.42	4.47		
NT2RP3004070	3.58	3.58	9.92	6.26	4.4	5.47		
NT2RP3004075	4.16	4.16	11.23	12.62	11.88	13.3		
NT2RP3004078	2.6	2.6	5.25	4.94	4.19	2.79		
NT2RP3004083	2.93	2.93	6.23	4.57	6.8	11.37		
NT2RP3004084	4.65	4.65	20.29	6.18	8.56	5.32		
NT2RP3004087	4.2	4.2	7.86	7.14	10.81	9.03		
NT2RP3004090	4.11	4.11	6.42	9	8.19	8.61	**	+
NT2RP3004093	2.38	2.38	7.49	4.07	3.51	4.47		
NT2RP3004095	5.02	5.02	13.11	11.57	10.17	18.55		
NT2RP3004102	3.32	3.32	5.59	5.25	4.27	3.21		
NT2RP3004110	12.74	12.74	18.66	22.12	14.31	19.97		
NT2RP3004119	3.3	3.3	7.71	3.91	4.08	3.73		
NT2RP3004125	5.55	5.55	12.05	8.13	10.88	8.38		
NT2RP3004129	4.62	4.62	7.38	3.36	2.95	6.08		
NT2RP3004130	11.81	11.81	28.12	21.92	31.13	21.05		
NT2RP3004133	4.51	4.51	12.95	14.62	8.94	16.81		
NT2RP3004145	1.43	1.43	4.17	2.62	4.87	3.59		
NT2RP3004148	2.67	2.67	7.07	5.26	6.24	4.5		
NT2RP3004155	2.37	2.37	4.82	4.7	4.57	6.59		
NT2RP3004165	17.94	17.94	29.96	29.58	31.82	39.18		
NT2RP3004179	7.34	7.34	6.72	2.71	5.41	3.3	*	-
NT2RP3004185	5.2	5.2	5.53	2.76	2.76	1.95	**	-
NT2RP3004188	4.77	4.77	10.82	7.74	11.35	7		
NT2RP3004189	4.23	4.23	5.91	4.97	4.82	6.28		
NT2RP3004190	2.6	2.6	5.57	5.84	4.36	5.26		
NT2RP3004191	14.09	14.09	23.4	31.41	29.45	30.09	*	+
NT2RP3004202	2.04	2.04	4.56	4.16	4.42	2.3		
NT2RP3004205	8.75	8.75	21.54	21.27	25.35	20.28		
NT2RP3004206	4.5	4.5	9.74	5.14	6.37	9		
NT2RP3004207	5.19	5.19	4.99	3.09	3.25	1.77	**	-
NT2RP3004209	4.74	4.74	7.74	8.2	11.23	9.08	*	+
NT2RP3004215	1.86	1.86	6.7	3.96	2.41	4.55		
NT2RP3004219	5.15	5.15	11.25	10.04	8.81	13.65		
NT2RP3004242	4.65	4.65	10.36	9.8	10.19	14.56		
NT2RP3004246	4.5	4.5	9.39	9.18	10.95	3.8		
NT2RP3004253	1.89	1.89	4.85	3.64	4.99	2.8		
NT2RP3004258	5.45	5.45	10.89	12.77	11.07	11.39		
NT2RP3004262	4.26	4.26	5.71	2.63	2.99	2.01	*	-
NT2RP3004275	5.59	5.59	3.43	1.4	2.97	2.34	*	-
NT2RP3004282	5.45	5.45	68.08	51.29	52.72	53.57		
NT2RP3004289	1.79	1.79	2.95	1.9	2.18	3.99		
NT2RP3004294	2.74	2.74	6.02	6.95	6.93	7.24	*	+
NT2RP3004298	8.76	8.76	48.63	46.33	60.89	50.83		
NT2RP3004309	3.3	3.3	6.46	5.2	5.22	5.31		
NT2RP3004321	3.71	3.71	6.11	3.29	3.74	3.34		

(1145)

特開 2002-191363

2287

2288

NT2RP3004322	5.61	5.61	6.86	6.06	6.43	6.56		
NT2RP3004332	11.69	11.69	100.11	78.54	102.41	76.72		
NT2RP3004334	1.49	1.49	6.97	8.56	6.06	9.06		
NT2RP3004336	2.11	2.11	6.24	6.02	4.44	5.63		
NT2RP3004338	3.09	3.09	8.41	10.22	9.52	16.47		
NT2RP3004341	1.81	1.81	4.56	6.13	5.17	9.13		
NT2RP3004345	4.1	4.1	8.68	9.3	9.63	8.48		
NT2RP3004348	5.06	5.06	11.25	13.04	10.79	12.54		
NT2RP3004349	5	5	7.5	4.89	7.75	5.76		
NT2RP3004355	5.57	5.57	7.09	7.55	7.07	7.18		
NT2RP3004356	5.76	5.76	21.51	11.29	15.14	15.56		
NT2RP3004360	3.4	3.4	5.26	6.01	5.32	7.85		
NT2RP3004361	2.6	2.6	6.26	7.67	7.3	8.87	*	+
NT2RP3004374	3.06	3.06	10.09	8.8	6.6	5.75		
NT2RP3004378	10.48	10.48	18.57	28.26	24.09	34.81	*	+
NT2RP3004399	3.88	3.88	5.77	3.53	3.17	9.06		
NT2RP3004405	4.07	4.07	6.77	3.03	5.52	3.93		
NT2RP3004406	5.36	5.36	6.23	5.19	6.03	6.12		
NT2RP3004411	5.93	5.93	13.28	8.08	6.39	9.51		
NT2RP3004424	1.53	1.53	2.43	3.27	1.81	2.83		
NT2RP3004428	3.03	3.03	5.36	5.07	3.82	4.09		
NT2RP3004432	3.3	3.3	3.52	3.61	3.11	4.38		
NT2RP3004434	3.42	3.42	8.41	7.28	9.09	7.99		
NT2RP3004446	3.29	3.29	4.6	3.29	4.1	2.63		
NT2RP3004451	3.2	3.2	6.01	3.89	3.38	2.48		
NT2RP3004454	2.96	2.96	4.16	2.69	3.5	2.5		
NT2RP3004466	3.5	3.5	7.89	5.25	3.85	5.61		
NT2RP3004470	7.42	7.42	24.53	18.4	16.35	24.72		
NT2RP3004472	2.49	2.49	4.4	3.97	3.84	3.88		
NT2RP3004475	1.71	1.71	5.52	2.72	5.93	3.9		
NT2RP3004480	14.12	14.12	17.04	18.94	10.5	18.82		
NT2RP3004481	5.42	5.42	11.37	5.04	7.37	12.39		
NT2RP3004490	2.66	2.66	8.45	3.92	7.03	8.25		
NT2RP3004496	4.8	4.8	14.38	7.22	9.3	11.08		
NT2RP3004498	6.39	6.39	21.39	16.86	15.11	18.92		
NT2RP3004503	2.78	2.78	9.34	4.85	6.23	5.88		
NT2RP3004504	3.91	3.91	11.09	6.05	10.67	11.52		
NT2RP3004505	17.38	17.38	28.56	37.35	26.72	37.82		
NT2RP3004507	1.57	1.57	6.6	2.52	5.52	1.68		
NT2RP3004519	4.9	4.9	7.73	4.93	8.36	8.75		
NT2RP3004524	10.04	10.04	29.21	26.88	26.43	29.62		
NT2RP3004527	3.03	3.03	3.22	1.29	2.23	2.08	*	-
NT2RP3004534	3.08	3.08	10.44	5.58	12	7.38		
NT2RP3004539	4.45	4.45	12.17	9.8	8.97	12		
NT2RP3004541	2.65	2.65	11.06	5.48	9.44	8.25		
NT2RP3004544	3.54	3.54	8.89	6.62	5.24	9.48		
NT2RP3004551	3.46	3.46	6.75	6.6	6.6	6.98		
NT2RP3004552	2.76	2.76	4.33	2.84	3.22	4.98		
NT2RP3004557	5.68	5.68	8.73	9.74	15.44	13.16	*	+
NT2RP3004561	1.96	1.96	3.77	2.55	4.46	4.1		

(1146)

特開2002-191363

2289

2290

NT2RP3004566	3.09	3.09	11.55	7.2	10.05	8.79		
NT2RP3004569	2.21	2.21	7.09	4.63	5.36	6.91		
NT2RP3004572	4.37	4.37	6.83	7.08	5.55	7.07		
NT2RP3004578	2.35	2.35	5.38	4.15	4.27	3.24		
NT2RP3004584	4.76	4.76	28.36	34.99	37.13	30.29		
NT2RP3004588	2.38	2.38	4.89	1.6	3.7	3.28		
NT2RP3004594	2.25	2.25	5.9	5.67	6.49	8.94		
NT2RP3004603	34.16	34.16	99.64	80.2	102.6	97.27		
NT2RP3004612	4.71	4.71	12.17	5.3	3.36	5.34		
NT2RP3004617	1.09	1.09	2.32	2.49	3.3	2.39		
NT2RP3004618	4.61	4.61	5.9	2.49	5.21	5.9		
NT2RP3004625	3.97	3.97	8.17	4.55	6.92	7.1		
NT2RP3004635	4.76	4.76	7.83	1.52	2.86	3.47		
NT2RP3004640	10.61	10.61	62.15	59.33	67.97	48.32		
NT2RP3004642	8.04	8.04	29.31	22.82	26.12	25.12		
NT2RP3004647	3.5	3.5	5.65	5.89	7.35	6.88	*	+
NT2RP3004652	1.76	1.76	10.37	4.2	3.71	4.34		
NT2RP3004669	2.01	2.01	5.36	4.01	5.33	3.46		
NT2RP3004670	5.04	5.04	10.58	12.4	9.19	14.23		
NT2RP4000008	45.17	45.17	71.24	49.77	32.43	48.77		
NT2RP4000018	11.64	11.64	14.61	11.69	14.8	14.87		
NT2RP4000023	6.96	6.96	8.91	4.86	7.38	5.98		
NT2RP4000025	16.2	16.2	22.16	26.22	29.89	24.7	*	+
NT2RP4000035	6.3	6.3	12.01	11.28	15.33	11.01		
NT2RP4000041	14.46	14.46	34.8	22.01	17.41	23.68		
NT2RP4000049	2.64	2.64	6.34	6.59	6.88	5.3		
NT2RP4000050	2.24	2.24	6.87	3.54	4.48	4.05		
NT2RP4000051	4.66	4.66	10.5	10.58	10.02	8.96		
NT2RP4000063	20.51	20.51	33.5	26.77	31	18.33		
NT2RP4000065	7.54	7.54	9.24	11.85	15.01	11.47	*	+
NT2RP4000070	6.63	6.63	5.29	3.79	4.54	3.84	*	-
NT2RP4000074	6.55	6.55	13.63	10.29	15.39	9.52		
NT2RP4000078	3.41	3.41	9.55	12.2	11.66	12.56	*	+
NT2RP4000080	3.52	3.52	7.01	5.06	6.84	6.81		
NT2RP4000099	128.86	128.86	236.22	149.44	161.1	211.56		
NT2RP4000102	3.55	3.55	6.48	5.27	4.97	5.54		
NT2RP4000103	2	2	5.47	2.57	2.85	3.96		
NT2RP4000108	4.66	4.66	7.91	9.33	10.73	10.61	*	+
NT2RP4000109	18.89	18.89	22.84	19.87	24.15	16.14		
NT2RP4000111	4.56	4.56	6.38	4.87	5.27	4.85		
NT2RP4000112	5.62	5.62	10.14	12.45	8.7	11.81		
NT2RP4000115	2.94	2.94	3.62	4.95	7.82	10.93		
NT2RP4000129	2.18	2.18	4.58	3.02	5.04	5.03		
NT2RP4000137	3.36	3.36	10.05	5.34	8.72	11.4		
NT2RP4000138	7.21	7.21	10.91	17.75	20.19	19.17	**	+
NT2RP4000141	3.25	3.25	6.1	5.22	4.9	3.57		
NT2RP4000147	6.21	6.21	4.49	4.27	5.2	3.47		
NT2RP4000150	5.96	5.96	6.93	7.33	10.41	7.06		
NT2RP4000151	2.82	2.82	7.82	5.69	6.24	5.05		
NT2RP4000157	73.27	73.27	222.87	169.53	97.5	173.8		

(1147)

特開 2002-191363

2291

2292

NT2RP4000159	2.02	2.02	5.03	3.38	5.03	2.92		
NT2RP4000163	5.21	5.21	8.74	10.89	9.18	6.63		
NT2RP4000167	3.26	3.26	4.35	4.32	3.22	3.69		
NT2RP4000171	5.72	5.72	7	5.72	7.45	5.03		
NT2RP4000175	62.48	62.48	94.56	144.06	214.73	147.88	*	+
NT2RP4000180	17.17	17.17	106.15	88.61	123.6	81.8		
NT2RP4000185	7.64	7.64	31.76	31.82	23.23	34.01		
NT2RP4000192	1.04	1.04	4.78	4.19	3.6	3.05		
NT2RP4000194	3.13	3.13	7.53	7.58	4.98	6.36		
NT2RP4000196	6.81	6.81	43.94	35.57	46.56	41.91		
NT2RP4000210	5.63	5.63	9.71	9.96	8.35	9.27		
NT2RP4000212	5.59	5.59	8.88	8.57	8.41	9.08		
NT2RP4000214	5.53	5.53	10.21	5.72	6.68	11.29		
NT2RP4000216	8.89	8.89	7.36	7	12.63	10.08		
NT2RP4000218	3.45	3.45	9.78	7.25	6.07	5.77		
NT2RP4000223	21.18	21.18	177.28	121.8	125.75	125.3		
NT2RP4000243	16.52	16.52	54.51	42.94	41.61	51.74		
NT2RP4000246	17.75	17.75	37.97	26.43	11.42	27.5		
NT2RP4000250	16.86	16.86	31.23	24.92	23.85	26.39		
NT2RP4000256	4.38	4.38	8.45	4.99	6.44	4.67		
NT2RP4000257	32.45	32.45	44.22	48.67	57.52	39.19		
NT2RP4000259	7.07	7.07	6.96	8.48	13.07	9.08		
NT2RP4000261	4.18	4.18	8.13	4.07	4.55	6.48		
NT2RP4000262	8.36	8.36	11.02	7.11	10.05	7.64		
NT2RP4000263	2.6	2.6	5.14	3.47	3.67	3.78		
NT2RP4000280	5.84	5.84	11.62	9.05	11.58	14.23		
NT2RP4000286	4.73	4.73	10.16	4.38	6.77	11.94		
NT2RP4000290	5.77	5.77	5.42	3.18	3.19	5.5		
NT2RP4000291	42.53	42.53	73.59	62.12	70.23	70.61		
NT2RP4000301	3.31	3.31	20.59	22.93	34.62	26.63	*	+
NT2RP4000312	7.76	7.76	45.27	39.01	43.59	45.92		
NT2RP4000321	3.73	3.73	8.16	7.88	7.14	9.75		
NT2RP4000323	1.44	1.44	4.26	2.27	2.55	2.12		
NT2RP4000324	7.77	7.77	16.76	8.33	10.2	11.08		
NT2RP4000334	20.97	20.97	77.78	63.81	71.76	68.24		
NT2RP4000343	2.48	2.48	5.54	1.57	3.16	2.38		
NT2RP4000348	3.4	3.4	10.81	8.38	6.75	13.2		
NT2RP4000349	1.78	1.78	0.83	0.37	0.72	2.48		
NT2RP4000355	3.98	3.98	14.84	7.18	9.13	9.04		
NT2RP4000356	8.3	8.3	22.64	9.22	11.44	21.13		
NT2RP4000360	3.04	3.04	6.98	5.06	5.43	4.73		
NT2RP4000367	2.18	2.18	3.72	2.04	2.58	2.83		
NT2RP4000370	4.21	4.21	7.51	3.62	7.12	5.99		
NT2RP4000373	3.33	3.33	5.18	3.19	5.23	2.95		
NT2RP4000376	4.2	4.2	5.5	4.85	5.53	6.5		
NT2RP4000381	1.97	1.97	6.46	4.31	7.01	5.59		
NT2RP4000388	85.82	85.82	204.63	128.93	93.38	116.03		
NT2RP4000390	12.16	12.16	94.1	78.97	76.6	82.02		
NT2RP4000393	8.66	8.66	9.77	9.09	5.99	10.79		
NT2RP4000398	5.51	5.51	26.52	22.08	28.3	26.61		

(1148)

特開2002-191363

2293

2294

NT2RP4000406	6.68	6.68	15.61	12.95	15.02	10.08		
NT2RP4000407	6.17	6.17	11.52	9.41	14.74	9.13		
NT2RP4000413	1.79	1.79	3.63	1.15	2.35	2.12		
NT2RP4000415	7.59	7.59	26.11	18.6	21.68	21.68		
NT2RP4000417	7.76	7.76	26.64	14.47	12.79	19.19		
NT2RP4000423	3.52	3.52	7.56	7	6.98	6.46		
NT2RP4000424	2.51	2.51	7.2	3.07	5.03	3.27		
NT2RP4000447	10.3	10.3	64.21	64.91	71.82	68.48		
NT2RP4000448	5.59	5.59	6.67	4.32	3.56	6.27		
NT2RP4000449	5.67	5.67	20.48	14.2	19.45	14.67		
NT2RP4000453	3.53	3.53	6.32	9.02	10.32	10.12	**	+
NT2RP4000455	2.64	2.64	3.98	1.81	1.75	2.4		
NT2RP4000456	12.57	12.57	21.7	14.7	9.86	13.58		
NT2RP4000457	1.54	1.54	7.12	3.9	7.55	2.98		
NT2RP4000461	5.7	5.7	9.84	7.65	6.41	4.6		
NT2RP4000462	11.76	11.76	15.32	11.86	11.53	17.37		
NT2RP4000463	10.2	10.2	52.59	50.66	69.3	48.36		
NT2RP4000471	6.36	6.36	10.74	5.74	6.23	4.98		
NT2RP4000472	3.97	3.97	4.41	1.27	1.27	1.66	**	-
NT2RP4000476	27.14	27.14	52.56	74.95	94.93	65.35	*	+
NT2RP4000480	11.56	11.56	29.27	19.08	9.95	26.66		
NT2RP4000481	2.29	2.29	4.76	3.73	4.16	4.33		
NT2RP4000483	1.38	1.38	7.59	7.58	7.85	6.26		
NT2RP4000487	1.54	1.54	4.9	2.26	3.17	0.91		
NT2RP4000496	2.16	2.16	4.98	2.95	4.07	3.65		
NT2RP4000497	17.15	17.15	22.33	34.9	44.46	29.9	*	+
NT2RP4000498	10.46	10.46	21.39	20.15	30.33	24.49		
NT2RP4000500	2.43	2.43	3.21	2.03	1.49	1.39	*	-
NT2RP4000507	5.63	5.63	9.02	12.49	10.62	17.51	*	+
NT2RP4000515	12.85	12.85	88.89	90.3	101.29	96.12		
NT2RP4000516	8.77	8.77	26.09	19.63	21.76	21.82		
NT2RP4000517	3.22	3.22	5.73	4.69	5.89	4.79		
NT2RP4000518	3.42	3.42	7.4	4.47	6.05	3.82		
NT2RP4000519	2.9	2.9	5.76	2.18	2.22	1.77		
NT2RP4000524	4.2	4.2	3.44	2.05	1.72	1.3	**	-
NT2RP4000528	3.67	3.67	3.06	3.01	3.27	8.01		
NT2RP4000537	35.4	35.4	62.6	36.23	30.93	44.52		
NT2RP4000541	2.04	2.04	2.45	3.34	4.33	3.09	*	+
NT2RP4000543	2.93	2.93	8.94	7.96	9.72	8.75		
NT2RP4000545	4.03	4.03	6.38	4.99	6.43	4.78		
NT2RP4000546	3.34	3.34	5.93	5.53	4.9	6.03		
NT2RP4000549	23.81	23.81	56.48	41.6	51.57	38.82		
NT2RP4000556	7.36	7.36	13.04	14.69	15.54	12.07		
NT2RP4000557	6.1	6.1	4.53	1.82	3.97	6.27		
NT2RP4000558	30.12	30.12	94.28	68.16	57.01	73.2		
NT2RP4000560	14.8	14.8	52.31	49.75	58.69	56.12		
NT2RP4000568	1.72	1.72	3.83	5.6	6.08	4.46	*	+
NT2RP4000583	11.61	11.61	20.2	23.65	14.97	20.56		
NT2RP4000585	3.04	3.04	4.14	3.12	2.55	3.24		
NT2RP4000588	8.65	8.65	12.77	14.58	16.96	13.55	*	+

(1149)

特開 2002-191363

2295

2296

NT2RP4000590	24.89	24.89	41.97	41.86	50.81	32.65
NT2RP4000599	4.29	4.29	3.24	2.44	2.23	3.59
NT2RP4000603	14.08	14.08	33.32	31.06	21.01	29.12
NT2RP4000607	2.41	2.41	10.04	4.45	7.87	15.35
NT2RP4000614	6.14	6.14	15.21	15.57	12.53	15.19
NT2RP4000634	6.61	6.61	11	7.78	9.84	10.31
NT2RP4000638	3.59	3.59	7.77	8.45	5.8	4.73
NT2RP4000648	3.13	3.13	4.26	2.54	2.69	2.19
NT2RP4000657	9.94	9.94	15.38	15.95	18.93	14.49
NT2RP4000691	5.76	5.76	5.82	4.92	7.47	7.73
NT2RP4000697	3.74	3.74	8.5	5.55	6.56	6.12
NT2RP4000704	8.91	8.91	47.2	44.17	54.81	38.14
NT2RP4000710	40.22	40.22	98.85	90.4	59.28	83.71
NT2RP4000713	4.35	4.35	19.92	16.67	20.85	15.52
NT2RP4000724	6.29	6.29	12.5	8.19	9.81	7.83
NT2RP4000725	3.61	3.61	4	1.88	1.74	2.33
NT2RP4000728	10.13	10.13	41.12	43.53	66.46	39.83
NT2RP4000737	4.07	4.07	2.15	3.63	3.09	3.28
NT2RP4000739	5.07	5.07	7.71	4.61	3.63	5.84
NT2RP4000749	2.4	2.4	5.29	2.59	3.97	1.68
NT2RP4000769	4.93	4.93	10.12	4.67	6.27	6.2
NT2RP4000774	3.34	3.34	8.87	5.12	6.63	4.27
NT2RP4000781	2.15	2.15	5.12	2.06	2.26	1.55
NT2RP4000783	6.81	6.81	15.16	13.48	15.44	12.67
NT2RP4000787	1.45	1.45	2.27	0.31	0.51	0.54
NT2RP4000788	3.58	3.58	23.26	16	18.3	18.73
NT2RP4000792	3.68	3.68	5.64	5.5	5.8	9.45
NT2RP4000809	43.7	43.7	56.09	46.75	50.47	81.62
NT2RP4000817	3.65	3.65	7.83	7.92	7.25	5.82
NT2RP4000821	31.34	31.34	38.66	28.32	33.11	25.22
NT2RP4000822	2.46	2.46	5.91	4.29	6.19	2.6
NT2RP4000823	697.74	697.74	1127.48	923.16	1026.8	947.85
NT2RP4000831	9.98	9.98	61.97	44.37	68.47	50.69
NT2RP4000833	3.19	3.19	11.26	6.73	7.19	11.91
NT2RP4000837	1.41	1.41	4.03	1.56	3.65	2.29
NT2RP4000839	12.23	12.23	97.13	79.71	85.74	86.06
NT2RP4000846	3.8	3.8	10.13	4.65	3.46	6.65
NT2RP4000848	4.63	4.63	10.74	8.65	8.58	6.07
NT2RP4000855	2.91	2.91	4.7	4	3.85	3.43
NT2RP4000863	3.08	3.08	4.33	3.11	5.3	3.78
NT2RP4000865	6.43	6.43	25.36	20.09	39.64	21.24
NT2RP4000873	9.64	9.64	88.25	63.22	69.65	71.33
NT2RP4000874	1.76	1.76	3.98	2.37	3.67	2.03
NT2RP4000875	3.31	3.31	9.24	6.88	6.52	7.19
NT2RP4000878	24.17	24.17	42.53	28.01	16.35	29.04
NT2RP4000879	2.56	2.56	5.1	2.95	5.29	2.62
NT2RP4000880	5.17	5.17	21.59	20.97	27.22	16.8
NT2RP4000891	81.07	81.07	192.57	252.29	351.53	221.08
NT2RP4000894	5.16	5.16	9.81	8.53	4.8	6.97
NT2RP4000898	0.86	0.86	2.74	1.88	2.14	1.64

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(1150)

特開2002-191363

2297

2298

NT2RP4000899	9.63	9.63	29.48	24.01	20.85	23.95		
NT2RP4000907	2.14	2.14	3.58	1.74	4.04	0.81		
NT2RP4000908	4.62	4.62	9.67	7.51	5.9	5.87		
NT2RP4000910	14.4	14.4	104.68	124.04	197.74	160.9	*	+
NT2RP4000918	2.85	2.85	4.76	4.73	4.26	5.35		
NT2RP4000925	3.9	3.9	5.53	2.81	3.15	1.86		
NT2RP4000927	1.99	1.99	2.5	0.46	1.08	0.6	**	-
NT2RP4000928	3.11	3.11	6.8	4.48	4.22	5.18		
NT2RP4000929	1.44	1.44	3.68	1.94	2.86	0.84		
NT2RP4000946	0.92	0.92	3.41	1.78	3.22	1.47		
NT2RP4000947	1.71	1.71	3.51	1.94	3.31	1.79		
NT2RP4000949	4.94	4.94	7.84	3.88	5.48	2.38		
NT2RP4000955	4.17	4.17	6.34	2.07	2.86	0.54	*	-
NT2RP4000959	20.55	20.55	28.14	36.21	42.82	36.14	*	+
NT2RP4000962	3.4	3.4	4.24	8.33	10.09	4.53		
NT2RP4000973	8.6	8.6	16.04	10.31	8.92	9.03		
NT2RP4000975	2.18	2.18	5.84	3.29	3.05	2.62		
NT2RP4000979	3.83	3.83	8.67	5.55	8.13	7.1		
NT2RP4000984	1.61	1.61	4.31	3.15	3.93	1.85		
NT2RP4000986	7.32	7.32	13.27	12.66	12.35	2.52		
NT2RP4000988	5.74	5.74	8.37	4	5.2	2.63		
NT2RP4000989	6.24	6.24	6.55	4.05	3.48	2.89	**	-
NT2RP4000990	3.16	3.16	4	1.92	1.69	2.16	**	-
NT2RP4000994	4.04	4.04	7.67	16.48	10.13	15.95	*	+
NT2RP4000996	3.54	3.54	7.49	6.77	6.52	7.38		
NT2RP4000997	21.59	21.59	36.81	28.52	15.18	34.38		
NT2RP4001001	5.53	5.53	9.17	16.66	18.38	15.09	**	+
NT2RP4001004	1.71	1.71	4.88	2.84	3.09	1.37		
NT2RP4001006	3.46	3.46	8.12	6.85	6.52	6.13		
NT2RP4001009	9.3	9.3	10.45	15.44	20.46	8.25		
NT2RP4001010	7.33	7.33	9.13	7.38	9.75	6.68		
NT2RP4001013	23.29	23.29	50.16	30.87	28.1	30.91		
NT2RP4001029	2.49	2.49	5.95	4.05	2.84	3.63		
NT2RP4001036	7.55	7.55	13.55	9.11	11.51	13.16		
NT2RP4001041	6.57	6.57	14.4	9.89	12.3	6.35		
NT2RP4001042	4.34	4.34	8.11	9.44	12.5	8.79		
NT2RP4001046	6.98	6.98	9.95	13.24	16.28	15.36	**	+
NT2RP4001050	5.28	5.28	4.81	3.79	4.64	3.35	*	-
NT2RP4001051	6.48	6.48	8.44	5.43	6.82	5.26		
NT2RP4001057	0.76	0.76	2.19	2.34	2.43	1.87		
NT2RP4001063	1.48	1.48	4.39	3.34	3.53	1.8		
NT2RP4001064	3.51	3.51	9.18	12.02	9.13	11.57		
NT2RP4001067	4.42	4.42	9.77	10.96	9.63	6.6		
NT2RP4001078	2.12	2.12	3.43	2.67	2.53	1.82		
NT2RP4001079	5.3	5.3	9.35	8.51	8.02	8.98		
NT2RP4001080	4.1	4.1	5.27	3.52	4.52	2.3		
NT2RP4001086	5.08	5.08	4.19	3.93	6.64	2.85		
NT2RP4001095	2.49	2.49	7.25	7.96	6.49	6.85		
NT2RP4001098	0.92	0.92	3.38	3.87	2.95	3.41		
NT2RP4001100	6.47	6.47	24.34	20.89	20.64	16.99		

(1151)

特開 2002-191363

2299

2300

NT2RP4001105	3.13	3.13	7.23	6.51	5.58	4.61		
NT2RP4001110	1.75	1.75	3.5	7.07	8.35	5.29	*	+
NT2RP4001115	9.95	9.95	17.68	20.6	18.48	15.31		
NT2RP4001117	19.81	19.81	30.49	35.35	42.53	27.5		
NT2RP4001122	6.06	6.06	6.09	5.17	6.25	3.27		
NT2RP4001123	3.62	3.62	7.76	7.95	5.96	6.27		
NT2RP4001126	4.36	4.36	11.28	10.87	9.09	8.04		
NT2RP4001127	3.25	3.25	4.59	3.39	3.08	2.17		
NT2RP4001138	2.46	2.46	5.8	3.41	2.56	1.62		
NT2RP4001143	2.73	2.73	5.98	6.44	6.54	5.66		
NT2RP4001148	3.72	3.72	6.76	3.77	3.03	2.05		
NT2RP4001149	5.07	5.07	7.28	6.76	9.03	6.37		
NT2RP4001150	3.8	3.8	3.17	3.15	3.7	2.88		
NT2RP4001159	7.08	7.08	11.61	7.69	5.58	10.46		
NT2RP4001162	3.77	3.77	6.14	4.07	6.06	3.41		
NT2RP4001170	1.15	1.15	4.53	1.28	3.2	2.15		
NT2RP4001174	4.16	4.16	12.27	7.91	11.95	5.02		
NT2RP4001175	9.65	9.65	19.14	15.72	21.29	13.28		
NT2RP4001176	99.19	99.19	161.51	174.03	241.92	194	*	+
NT2RP4001184	4.83	4.83	27.1	25.76	32.51	22.85		
NT2RP4001198	21.66	21.66	48.22	29.54	29.17	38.54		
NT2RP4001199	2.52	2.52	6.45	2.59	4.74	3.71		
NT2RP4001206	8.25	8.25	33.2	25.92	32.07	25.48		
NT2RP4001207	2.38	2.38	5.15	2.21	3.11	4.01		
NT2RP4001210	2.73	2.73	5.2	3.62	4.26	2.64		
NT2RP4001213	3.42	3.42	5.11	3.99	4.23	3.63		
NT2RP4001214	3.34	3.34	4.3	3.51	3.76	2.16		
NT2RP4001219	7.4	7.4	12.05	14.35	19.28	13.39	*	+
NT2RP4001228	5.26	5.26	9.63	12.15	15.74	20.07	*	+
NT2RP4001235	2.42	2.42	7.45	3.46	6.02	4.48		
NT2RP4001256	2.11	2.11	4.24	1.66	3.41	2.66		
NT2RP4001257	2.48	2.48	7.27	4.05	4.35	4.05		
NT2RP4001260	3.16	3.16	5.79	2.52	3.86	2.92		
NT2RP4001261	3.84	3.84	6.63	8.42	6.47	5		
NT2RP4001274	22.92	22.92	38.08	25.02	31.56	21.25		
NT2RP4001276	5.24	5.24	10.03	11.38	15.97	11.63	*	+
NT2RP4001283	20.72	20.72	122.55	87.44	93.43	86.47		
NT2RP4001299	9.62	9.62	15.14	14.95	10.52	15.18		
NT2RP4001313	1.45	1.45	3.26	1.72	2.96	0.97		
NT2RP4001315	6.06	6.06	11.14	7.45	9.92	7.74		
NT2RP4001320	14.6	14.6	42.74	32.02	38.13	29.24		
NT2RP4001325	32.53	32.53	146.14	142.88	178.36	128.89		
NT2RP4001336	6.69	6.69	40.75	38.55	46.66	32.11		
NT2RP4001339	4.12	4.12	5.6	3.35	5.56	2.76		
NT2RP4001343	10.46	10.46	83.37	54.71	61.01	60.06		
NT2RP4001344	6.7	6.7	60.08	49.79	55.21	42.62		
NT2RP4001345	1.65	1.65	6.68	5.64	5.7	3.87		
NT2RP4001351	4.1	4.1	15.97	10.01	20.05	11.42		
NT2RP4001353	2.8	2.8	5.91	1.63	2.94	1.86		
NT2RP4001355	2.57	2.57	8.67	1.83	3.12	2.08		

(1152)

特開 2002-191363

2301

2302

NT2RP4001367	10.64	10.64	17.66	11.92	17.06	13.16		
NT2RP4001372	2.26	2.26	3.82	2.1	2.06	2.07		
NT2RP4001373	8.86	8.86	16.4	10.99	8.59	11.48		
NT2RP4001375	2.71	2.71	6.06	4.91	7.42	2.94		
NT2RP4001379	1.74	1.74	3.52	2.34	4.67	1.38		
NT2RP4001381	5.6	5.6	12.51	12.8	12.66	8.27		
NT2RP4001386	6.39	6.39	14.52	14.77	20.11	12.32		
NT2RP4001389	7.28	7.28	8.66	5.43	9.25	4.09		
NT2RP4001396	5.76	5.76	6.42	2.83	4.61	2.12	*	-
NT2RP4001407	2.92	2.92	2.98	3.07	2.04	1.76		
NT2RP4001409	13.6	13.6	28.28	8.87	5.85	8.84		
NT2RP4001410	33.56	33.56	61.26	40.57	18.92	37.8		
NT2RP4001414	16.59	16.59	37.89	14.29	21.3	16.8		
NT2RP4001424	3.55	3.55	8.85	7.99	10.45	7.41		
NT2RP4001433	3.85	3.85	6	8.39	9.79	3.38		
NT2RP4001438	9.95	9.95	27.94	46.22	53.63	30.76	*	+
NT2RP4001442	4.33	4.33	4.97	3.3	3.41	2.64	*	-
NT2RP4001447	4.42	4.42	4.69	5.08	5.51	3.41		
NT2RP4001466	3.74	3.74	5.45	7.38	3.23	5.57		
NT2RP4001467	21.67	21.67	58.89	54.18	44.07	55.8		
NT2RP4001472	8.05	8.05	12.19	11.7	11.76	11.97		
NT2RP4001474	2.83	2.83	4.81	3.2	5.73	1.98		
NT2RP4001483	1.48	1.48	4.33	2.61	3.7	1.19		
NT2RP4001488	21.03	21.03	32.41	39.07	46.9	33.42		
NT2RP4001492	4.18	4.18	6.73	3.21	3.81	1.91		
NT2RP4001498	4.3	4.3	2.43	2.95	3.52	2.13		
NT2RP4001502	28.2	28.2	57.38	27.65	34.81	41.83		
NT2RP4001503	3.83	3.83	6.74	5.51	4.1	3.37		
NT2RP4001507	2.39	2.39	3.69	5.03	5.95	5.18	**	+
NT2RP4001510	1.74	1.74	4.63	5.64	5.05	3.02		
NT2RP4001516	3.54	3.54	4.16	2.52	1.9	1.27	*	-
NT2RP4001520	25.33	25.33	70.64	73.93	107.21	85.05		
NT2RP4001523	5.57	5.57	9.99	8.4	7.79	6.19		
NT2RP4001524	6.1	6.1	10.17	8.92	11.45	5.98		
NT2RP4001529	29.8	29.8	74.89	59.39	60.69	55.29		
NT2RP4001531	2.88	2.88	10.96	8.63	11.05	7.81		
NT2RP4001546	143.48	143.48	388.31	316.63	215.6	309.45		
NT2RP4001547	9.05	9.05	26.31	28.76	23.06	18.81		
NT2RP4001551	2.27	2.27	3.93	3.87	4.08	2.02		
NT2RP4001555	2.95	2.95	5.8	4.14	3.58	1.74		
NT2RP4001567	6.54	6.54	6.88	5.82	7.12	5.68		
NT2RP4001568	7.58	7.58	11.65	10.02	19.33	12.97		
NT2RP4001569	15.49	15.49	41.42	32.56	40.11	27.18		
NT2RP4001571	5.75	5.75	13.28	9.06	7.43	8.13		
NT2RP4001574	8.5	8.5	19.03	15.54	17.52	15.29		
NT2RP4001575	2.46	2.46	5.86	5.71	5.53	4.24		
NT2RP4001578	17.21	17.21	45.64	52.76	53.63	40.92		
NT2RP4001592	9.76	9.76	13.68	14.13	19.64	11.68		
NT2RP4001593	9.4	9.4	18.11	22.12	27.3	18.04		
NT2RP4001605	5.97	5.97	4.78	4.1	7.77	4.66		

(1153)

特開2002-191363

2303

2304

NT2RP4001606	2.9	2.9	8.34	6.01	3.51	6.75		
NT2RP4001607	2.04	2.04	5.24	4.57	3.26	3.78		
NT2RP4001610	1.74	1.74	2.42	2.6	2.48	1.7		
NT2RP4001614	2.17	2.17	7.19	5.38	4.34	6.86		
NT2RP4001623	2.38	2.38	5.26	2.43	2.65	2.02		
NT2RP4001626	9.48	9.48	11.67	18.67	23.9	19.44	**	+
NT2RP4001634	2.74	2.74	4.93	3.67	5.24	4.26		
NT2RP4001638	3.41	3.41	3.03	2.36	2.11	1.87	**	-
NT2RP4001644	7.86	7.86	33.73	24.36	26.04	24.99		
NT2RP4001646	11.61	11.61	15.02	7.42	10.74	11.21		
NT2RP4001656	3.75	3.75	5.23	2.89	4.51	2.07		
NT2RP4001666	1.99	1.99	4.68	3.26	6.25	2.02		
NT2RP4001670	11.74	11.74	15.51	12.45	7.09	8.31		
NT2RP4001677	28.27	28.27	42.75	42.01	45.48	47.53		
NT2RP4001679	8.82	8.82	33.83	33.23	51.5	29.78		
NT2RP4001695	7.71	7.71	12.76	15.66	20.35	12.87		
NT2RP4001696	2.64	2.64	5.45	3.13	3.92	3.72		
NT2RP4001699	3.58	3.58	8.03	3.18	4.12	4.42		
NT2RP4001717	2.79	2.79	7.03	3.29	5.84	4.15		
NT2RP4001719	3.59	3.59	9.11	7.6	9.27	6.28		
NT2RP4001725	2.27	2.27	4.79	2.28	5.07	1.43		
NT2RP4001726	7.07	7.07	11.18	5.85	6.91	4.98		
NT2RP4001730	3.11	3.11	12.82	11.96	19.81	16.3		
NT2RP4001739	2.83	2.83	5.83	5.79	6.55	4.98		
NT2RP4001741	7.25	7.25	15.93	9.28	12.42	10.97		
NT2RP4001753	3.04	3.04	8.4	4.39	4.64	6.64		
NT2RP4001760	4.32	4.32	6.6	7.79	7.73	12.96		
NT2RP4001787	67.61	67.61	173.05	169.17	187.1	193.22		
NT2RP4001790	2	2	5.29	3.42	2.97	2.58		
NT2RP4001795	9.31	9.31	12.31	14.38	19.76	12.34		
NT2RP4001803	3.35	3.35	3.6	3.94	4.78	3.67		
NT2RP4001805	2.64	2.64	3.57	3.64	2.47	2.95		
NT2RP4001809	4.84	4.84	26.35	18.18	23.17	11.33		
NT2RP4001817	11.55	11.55	19.09	9.5	10.78	12.71		
NT2RP4001822	2.09	2.09	5.36	3.73	5.11	3.33		
NT2RP4001823	1.91	1.91	3.95	1.14	3.34	1.24		
NT2RP4001827	14.88	14.88	25.96	35.78	40.37	29.5	*	+
NT2RP4001828	9.76	9.76	34.37	27.78	34.21	26.3		
NT2RP4001836	7.74	7.74	33.26	27.19	39.14	26.78		
NT2RP4001838	1.59	1.59	7.49	2.09	4.5	2.71		
NT2RP4001841	8.75	8.75	80.37	61.67	50.27	56.46		
NT2RP4001849	1.9	1.9	4.55	2.51	5.08	1.58		
NT2RP4001861	7.27	7.27	34.6	36.09	41.9	34.39		
NT2RP4001877	6.59	6.59	8.44	12.87	9.04	14.1	*	+
NT2RP4001879	9.64	9.64	15.47	11.58	14.24	10.73		
NT2RP4001889	5.09	5.09	10.66	6.68	11.25	8.91		
NT2RP4001893	3.97	3.97	7.34	3.19	6.11	2.72		
NT2RP4001896	3.18	3.18	6.86	5.38	7.87	4.92		
NT2RP4001898	7.83	7.83	26.41	22.98	20.13	20.15		
NT2RP4001901	1.73	1.73	4.69	4.08	5.87	2.69		

(1154)

特開 2002-191363

2305

2306

NT2RP4001910	39.51	39.51	58.21	53.45	33.59	45.93		
NT2RP4001925	4.1	4.1	10.21	6.69	6.32	6.12		
NT2RP4001926	6.41	6.41	7.22	7.1	9.54	5.52		
NT2RP4001927	6.26	6.26	9.97	5.83	8.13	2.82		
NT2RP4001931	8.64	8.64	14.16	15.49	17.54	11.89		
NT2RP4001933	38.49	38.49	87.13	96.81	133.51	91.22		
NT2RP4001938	2.93	2.93	4.53	4.27	3.31	3.55		
NT2RP4001942	13.44	13.44	27.12	31.34	24.8	18.71		
NT2RP4001945	1.41	1.41	4	1.55	3.67	1.77		
NT2RP4001946	1.97	1.97	5.67	3.04	3.96	1.66		
NT2RP4001947	4.42	4.42	8.93	5.92	9.81	5.16		
NT2RP4001950	4.13	4.13	5	2.25	2.84	1.34	*	-
NT2RP4001953	10.44	10.44	14.15	13.81	19.4	14.36		
NT2RP4001966	2.44	2.44	2.41	2.51	4.26	1.52		
NT2RP4001970	2.26	2.26	5.32	3.91	3.4	2.88		
NT2RP4001975	8.56	8.56	20.03	18.32	13.05	12.02		
NT2RP4001988	6.72	6.72	18.78	22.92	24.78	29.44	*	+
NT2RP4001996	5.27	5.27	12.83	10.65	16.35	12.42		
NT2RP4002014	3.4	3.4	8.14	43.19	37.87	33.17	**	+
NT2RP4002018	6.19	6.19	13.71	10.47	11.39	10.36		
NT2RP4002035	5.35	5.35	5.95	5.4	4.17	2.54		
NT2RP4002043	7.1	7.1	10.8	9.64	12.2	6.5		
NT2RP4002046	9.74	9.74	20.08	21.94	15.28	17.11		
NT2RP4002047	8.37	8.37	19.18	22.28	24.07	28.83	*	+
NT2RP4002052	5.78	5.78	10.36	9.02	9.36	9.37		
NT2RP4002056	32.58	32.58	71.49	58.09	76.58	49.75		
NT2RP4002057	6.37	6.37	11.06	12.13	13.58	7.41		
NT2RP4002058	3.85	3.85	6.6	4.1	4.2	3.22		
NT2RP4002064	5.93	5.93	4.39	2.6	4.16	2.07	*	-
NT2RP4002071	6.67	6.67	7.07	6.95	10.06	6.27		
NT2RP4002075	1.16	1.16	2.11	2.27	2.35	1.27		
NT2RP4002078	2.25	2.25	8.63	6.86	8.24	4.97		
NT2RP4002081	8.11	8.11	26.15	18.73	18.78	19.42		
NT2RP4002083	1.39	1.39	5.25	3.36	3.16	1.88		
NT2RP4002099	3.26	3.26	4.73	2.84	3.56	2.51		
NT2RP4002106	10.35	10.35	20.34	22.36	25.93	20.55		
NT2RP4002111	11.7	11.7	12.37	19.77	30.44	17.72		
NT2RP4002112	6.15	6.15	10.97	8.9	8.34	3.22		
NT2RP4002116	12.6	12.6	47.19	37.43	41.25	28.65		
NT2RP4002122	5.34	5.34	9.29	14.84	14.86	12.67	**	+
NT2RP4002126	6.42	6.42	14.44	16.82	14.35	10.42		
NT2RP4002133	7.56	7.56	20.82	29.17	26.14	21		
NT2RP4002136	3.63	3.63	5.74	4.89	5.38	2.69		
NT2RP4002139	26.89	26.89	31.12	60.65	61.92	32.88		
NT2RP4002174	139.27	139.27	232.64	240.71	275.01	193.19		
NT2RP4002185	7.77	7.77	13.2	12.36	19.06	11.58		
NT2RP4002186	4.5	4.5	9.83	7.82	4.72	6.78		
NT2RP4002187	15.42	15.42	32.13	26.94	19.84	21.17		
NT2RP4002188	3.01	3.01	8.34	8.3	7.75	6.41		
NT2RP4002199	1.85	1.85	3.73	2.6	2.91	3.78		

(1155)

特開2002-191363

2307

2308

NT2RP4002206	2.08	2.08	3.39	2.48	2.34	1.29
NT2RP4002210	3.13	3.13	4.75	2.02	2.86	0.98
NT2RP4002222	4.2	4.2	6.63	5.56	6.28	4.16
NT2RP4002241	7.97	7.97	8.24	10.82	11.75	7.19
NT2RP4002248	5.08	5.08	16.69	10.74	8.9	8.13
NT2RP4002250	1.54	1.54	3.22	0.73	1.69	0.56
NT2RP4002259	4.86	4.86	9.82	3.21	4.85	1.75
NT2RP4002268	16.62	16.62	29.54	28.9	28.18	25.68
NT2RP4002288	6.42	6.42	12.57	13.29	14.36	11.97
NT2RP4002290	7.55	7.55	7.61	7.96	7.67	5.87
NT2RP4002298	3.92	3.92	4.18	5.54	5.03	4.18
NT2RP4002306	2.38	2.38	5.79	2.97	5.77	2.64
NT2RP4002308	2.04	2.04	6.03	5.31	5.23	4.1
NT2RP4002336	2.71	2.71	6.33	3.71	4.19	4.63
NT2RP4002340	1.09	1.09	3.96	1.28	2.75	0.49
NT2RP4002361	2.77	2.77	5.78	3.73	4.03	2.48
NT2RP4002367	2.27	2.27	5.84	3.23	2.48	2.77
NT2RP4002368	9.87	9.87	17.2	18.26	19.27	16
NT2RP4002377	3.3	3.3	23.8	25.46	30.75	23.93
NT2RP4002408	2.22	2.22	3.87	3.75	6.37	4.11
NT2RP4002425	2.84	2.84	5.81	8.24	7.98	5.23
NT2RP4002432	12.33	12.33	85.4	61.06	72.53	67.82
NT2RP4002447	2.97	2.97	7.68	3.96	5.4	4.59
NT2RP4002451	5.48	5.48	6.2	5.84	5.85	6.83
NT2RP4002461	9.8	9.8	32.09	32.76	38.91	29.04
NT2RP4002486	3.5	3.5	6.71	2.47	4.15	2.87
NT2RP4002517	3.65	3.65	9.11	7.02	8.53	7.18
NT2RP4002556	4.29	4.29	3.91	5.68	10.03	6.41
NT2RP4002569	3.36	3.36	7.36	4.93	5.29	3.42
NT2RP4002587	2.26	2.26	4.19	2.8	3.4	2.02
NT2RP4002591	2.21	2.21	4.89	2.89	4.5	3.08
NT2RP4002607	1.43	1.43	3.34	2.87	4.63	1.58
NT2RP4002627	17.83	17.83	61.9	55.9	76.17	65.3
NT2RP4002628	7.28	7.28	15.48	14.53	23.95	12.54
NT2RP4002630	4.19	4.19	5.25	6.72	9.4	7.16
NT2RP4002639	9.43	9.43	70.25	52.38	77.24	57.28
NT2RP4002641	1.58	1.58	9.03	3.94	4.07	4.1
NT2RP4002658	114.62	114.62	166.93	76.49	34.96	109.83
NT2RP4002669	3.5	3.5	5.67	5.4	5.33	4.68
NT2RP4002677	6.24	6.24	9.41	10.14	7.99	13.62
NT2RP4002715	8.42	8.42	34.92	40.1	48.46	32.3
NT2RP4002750	2.6	2.6	8.29	1.68	2.04	1.33
NT2RP4002784	3.71	3.71	9.51	9.44	11.22	7.06
NT2RP4002791	4.91	4.91	9.44	4.88	9.76	5.33
NT2RP4002811	1.63	1.63	6.38	3.17	2.95	3.43
NT2RP4002830	4.26	4.26	7.45	3.9	5.9	5.46
NT2RP4002832	2.12	2.12	3.13	2.38	5.59	2.54
NT2RP4002850	5.07	5.07	12.04	14.36	12.63	8.06
NT2RP4002874	5.17	5.17	6.67	3.41	5.14	1.96
NT2RP4002884	28.81	28.81	49.75	43.57	74.75	52.87

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(1156)

特開2002-191363

2309

2310

NT2RP4002888	5.55	5.55	4.83	3.67	4.32	3.08	*	-
NT2RP4002891	5.48	5.48	15.79	13.16	19.42	11.91		
NT2RP4002894	12.04	12.04	24.47	18.44	12.76	16.4		
NT2RP4002896	5.54	5.54	12.2	8.96	6.18	7.78		
NT2RP4002905	1.71	1.71	4.27	2.32	3.58	1.28		
NT2RP4002907	5.11	5.11	7.62	6.94	10.72	1.41		
NT2RP5003459	68.11	68.11	133.25	154.61	146.15	164.37	*	+
NT2RP5003461	7.34	7.34	10.14	10.85	14.36	8		
NT2RP5003471	106.6	106.6	168.71	124.4	148.85	112.14		
NT2RP5003477	2.71	2.71	2.62	2.59	2.33	1.9		
NT2RP5003487	157.44	157.44	424.89	292.71	256.56	354.93		
NT2RP5003492	3.1	3.1	4.91	5.25	6.17	5.91	*	+
NT2RP5003500	1.5	1.5	3.28	2.38	2.54	2.59		
NT2RP5003506	4.96	4.96	9.3	7.83	10.37	9.04		
NT2RP5003512	2.21	2.21	4.35	2.63	3.46	2.15		
NT2RP5003522	4.1	4.1	5.97	4.62	4.19	2.34		
NT2RP5003524	4.38	4.38	3.86	1.61	1.54	0.84	**	-
NT2RP5003527	24.72	24.72	71.27	76.81	87.24	60.59		
NT2RP5003531	7.16	7.16	17.2	15.58	14.06	14.11		
NT2RP5003534	2.68	2.68	5.49	5.54	6.82	4.55		
NT2RP6000020	8.69	8.69	19.96	14.65	15.13	16.29		
NT2RP6000022	3.19	3.19	4.05	4.06	3.96	2.44		
NT2RP6000050	3.95	3.95	3.99	4.98	5.82	2.88		
NT2RP6000063	3.91	3.91	6.04	3.61	2.52	2.56		
NT2RP6000074	5.38	5.38	4.88	3.41	3.27	2.17	**	-
NT2RP6000083	7.76	7.76	11.18	11.49	16	9.91		
NT2RP6000100	2.49	2.49	4.58	4.04	4.71	3.3		
NT2RP6000123	1.94	1.94	3.29	5.1	4.26	4.22	*	+
NT2RP6000129	1.9	1.9	4.47	4.06	4.27	2.74		
NT2RP6000147	3.75	3.75	11.74	10.8	11.03	7.48		
NT2RP6000163	2.62	2.62	4.23	2.28	1.95	1.71		
NT2RP6000181	8.03	8.03	12.4	9.44	13.25	9.01		
NT2RP6000182	5.44	5.44	6.42	4.82	5.56	3.88		
OVARC1000001	4.97	4.97	5.24	6.04	7.48	2.35		
OVARC1000003	3.21	3.21	8.31	8.51	7.66	7.05		
OVARC1000004	9.87	9.87	116.19	88.04	109.99	85.44		
OVARC1000006	3.57	3.57	6.58	9.04	7.88	4.73		
OVARC1000013	6.51	6.51	9.19	7.32	8.36	8.33		
OVARC1000014	3.39	3.39	5.02	4.23	5.02	4.17		
OVARC1000017	3.11	3.11	6.81	4.2	4.45	2.72		
OVARC1000026	24.79	24.79	32.1	56.82	69.34	44.53	*	+
OVARC1000035	11.11	11.11	20.26	20.41	23.65	15.36		
OVARC1000037	8.73	8.73	19.12	15.64	9.13	15.9		
OVARC1000058	6.06	6.06	11.69	13.84	7.56	10.6		
OVARC1000060	1.89	1.89	6.28	5.98	5.24	5.13		
OVARC1000068	2.38	2.38	5.33	5.31	3.56	2.42		
OVARC1000069	74.66	74.66	101.53	75.95	84.36	86.42		
OVARC1000071	4.4	4.4	4.77	6.47	5.35	4.04		
OVARC1000075	55.43	55.43	125.63	120.89	150.97	117.03		
OVARC1000083	9.58	9.58	9.24	13.12	12.7	10.64	*	+

(1157)

特開2002-191363

2311

2312

OVARC1000085	106.6	90.9	156.14	214.2	177.05	273.14	*	+
OVARC1000086	3.98	6.82	9.23	11.98	11.3	14.09	*	+
OVARC1000087	1.51	2.83	1.79	4.03	3.57	3.35	*	+
OVARC1000090	1.48	4.1	6.14	10.88	9.58	8.79	*	+
OVARC1000091	4.88	8.33	8.01	7.99	7.76	6.82		
OVARC1000092	2.83	6.81	4.18	4.68	6.25	4.85		
OVARC1000105	9.73	14.86	17.21	26.29	25.62	22.88	*	+
OVARC1000106	26.02	23.03	46.38	66.36	50.1	53.01	*	+
OVARC1000109	9.12	13.08	18.04	16.72	12.91	17.46		
OVARC1000113	4.12	6.25	6.53	6.83	8.19	7.65		
OVARC1000114	2.14	3.44	5.77	5.94	5.86	4.98		
OVARC1000133	2.53	4.96	6.36	4.05	4.97	2.95		
OVARC1000137	6.14	10.05	13.51	13.3	18.59	14.39		
OVARC1000139	14.75	20.77	83.44	71.14	98.1	69.29		
OVARC1000145	0.72	6.64	2.89	1.78	2.42	2		
OVARC1000148	5.09	4.98	7.88	4.91	5.32	7.91		
OVARC1000151	1.41	2.11	2.4	3.58	4.08	3.58	**	+
OVARC1000157	10.99	14.16	17.51	21.21	25.06	22.76	*	+
OVARC1000162	1.22	4.4	2.5	2.93	2.49	2.59		
OVARC1000168	1.98	8.46	6.2	8.01	9.61	9.96		
OVARC1000169	32.03	45.07	49.48	70.63	69.6	89.08	*	+
OVARC1000178	0.84	5.08	2.53	3.37	3.18	2.78		
OVARC1000182	0.8	3.3	1.42	2.02	1.95	1.78		
OVARC1000186	2.51	3.72	3.23	5.95	3.27	4.77		
OVARC1000188	1.04	2.67	2.33	2.48	2.87	1.9		
OVARC1000191	1.01	3.8	2.63	3.12	2.85	2.54		
OVARC1000198	2.09	3.59	4.32	5.62	5.12	5.06	*	+
OVARC1000208	6.49	10.37	22.5	17.79	24.54	22.02		
OVARC1000209	7.99	13.69	22.82	23.42	27.81	29.16		
OVARC1000212	2.47	5.63	3.59	4.76	5.03	4.88		
OVARC1000216	1.72	4.96	4.36	15.43	11.3	12.54	**	+
OVARC1000240	2.98	3.53	8.13	5.39	5.46	4.87		
OVARC1000241	1.29	2.47	3.18	2.65	3.17	1.4		
OVARC1000249	4.14	5.43	8.17	5.46	5	6.13		
OVARC1000254	33.15	39.39	100.99	100.41	131.42	100.89		
OVARC1000255	0.85	4.83	2.51	2.98	2.45	1.95		
OVARC1000267	2.37	6.41	6.71	6.66	7.16	7.31		
OVARC1000275	79.02	93.7	161.08	199.43	240.76	175.96	*	+
OVARC1000287	226.67	224.66	236.08	433.91	512.76	470.31	**	+
OVARC1000288	3.2	4.25	7.38	6.23	5.32	4.47		
OVARC1000298	8.96	10.09	19.62	13.37	7.19	9.6		
OVARC1000302	1.12	2.14	2.13	2.47	1.85	2		
OVARC1000304	1.09	2.68	3.23	5.02	3.41	6		
OVARC1000307	2.95	6.19	4.74	7.59	4.7	6.29		
OVARC1000309	1.18	7.16	3.22	3.24	2.85	2.4		
OVARC1000312	2.83	11.64	6.03	4.17	5.4	2.46		
OVARC1000313	10.48	19.25	14.81	9.39	17.54	22.17		
OVARC1000321	31.6	24.05	47.79	30.5	31.37	15.43		
OVARC1000326	1.52	2.3	3.9	3.84	3.17	2.79		
OVARC1000327	1.52	3.28	4.24	3.13	1.49	2.46		

(1158)

特開2002-191363

2313

2314

OVARC1000331	2.22	4.72	2.41	4.33	4.45	4.58		
OVARC1000335	2.3	5.84	4.02	2.72	5.16	4.75		
OVARC1000347	1.83	8.18	6.24	7.35	9.24	8.44		
OVARC1000348	1.61	10.62	3.73	2.84	4.59	3.05		
OVARC1000363	3.7	9.61	6.51	7.31	11.52	6.83		
OVARC1000377	1.07	2.09	2.43	2.28	2.51	2.45		
OVARC1000382	3.34	3.39	4.33	5.07	2.52	1.03		
OVARC1000384	4.2	5.42	8.35	5.32	4.4	6.04		
OVARC1000401	0.62	3.63	2.09	3.35	4.08	3.64		
OVARC1000406	18.98	23.3	49.12	57.09	74.48	54.63	*	+
OVARC1000407	1.99	6.28	3.99	4.11	6.42	3.16		
OVARC1000408	27.5	38.45	70.39	74.84	111.17	71.25		
OVARC1000410	6.83	12.72	10.41	4.78	6.65	5.01		
OVARC1000411	0.91	1.5	2.6	3.49	3	2.22		
OVARC1000414	1.31	2.22	3.7	4.5	3.78	3.71		
OVARC1000420	1.44	2.76	3.29	3.3	2.59	2.1		
OVARC1000421	1.42	2.65	3.33	3.96	4.21	4.99	*	+
OVARC1000427	25.78	27.02	130.06	156.9	215.67	142.19		
OVARC1000431	10.51	17.6	19.12	33.66	31.78	25.78	*	+
OVARC1000437	3.14	6.37	7.31	5.97	7.63	6.36		
OVARC1000439	5.81	10.95	13.82	21.81	23.01	21.52	**	+
OVARC1000440	2.56	3.74	5.01	7.47	9.31	7.08	*	+
OVARC1000442	2.34	2.38	6.81	6.66	9.5	8.12		
OVARC1000443	2.09	2.2	2.88	3.29	3.41	2.62		
OVARC1000461	1.11	2.84	2.2	2.55	1.12	2.14		
OVARC1000465	3.27	5.01	3.51	3.94	4.62	3.95		
OVARC1000466	1.94	5.47	5.9	6.54	10.13	6.76		
OVARC1000467	1.01	5.08	2.41	3.65	2.98	3.78		
OVARC1000470	1.13	5.81	3.03	3.18	4.02	3.78		
OVARC1000473	1.81	1.95	2.65	2.44	4.16	1.39		
OVARC1000479	5.67	5.88	9.88	10.35	14.26	6.88		
OVARC1000484	3.99	5.74	6.54	8.66	9.93	6.87		
OVARC1000486	3.17	4.71	4.49	5.74	4.93	4.28		
OVARC1000496	0.93	3.55	0.66	0.31	1.07	0.62		
OVARC1000520	0.84	5.89	1.18	1.32	2.27	2		
OVARC1000522	4.1	7.19	12	13.85	14.03	10.34		
OVARC1000526	1.96	7.04	3.75	5.93	5.48	4.69		
OVARC1000529	2.38	2.57	4.44	3.66	4.16	3.08		
OVARC1000533	3.3	4.66	7.95	8.89	12.29	7.83		
OVARC1000543	0.84	2.44	2.06	2.57	3.08	2.72		
OVARC1000550	0.75	3.68	2.32	2.82	4.04	2.34		
OVARC1000553	2.1	5.63	7.02	7.72	7.69	6.95		
OVARC1000556	5.77	15.21	11.77	8.95	13.91	8.96		
OVARC1000557	0.83	5.12	1.4	1.61	2.29	1.88		
OVARC1000561	3.48	7.38	9.26	13	17.66	15.09	*	+
OVARC1000564	8.89	9.02	10.44	17.84	11.31	16.69	*	+
OVARC1000573	1.87	3.68	4.86	5.75	5.32	3.83		
OVARC1000576	24.12	29.23	124.94	83.09	93.83	98.58		
OVARC1000578	2.43	4.6	5.53	8.64	4.46	3.93		
OVARC1000581	0.34	3.28	1.15	1.75	1.27	1.23		

(1159)

特開2002-191363

2315

2316

OVARC1000586	22.54	28.9	41.17	34.58	43.39	40.93		
OVARC1000588	0.74	5.23	2.03	2.75	3.72	2.05		
OVARC1000605	1.98	2.62	2.88	4.47	4.23	3.87	**	+
OVARC1000622	3.86	4.59	11.57	12.7	11.13	11.6		
OVARC1000636	1.64	3.79	4.58	4.19	4.03	5.09		
OVARC1000640	1.97	4.72	3.93	4.21	3.67	3.09		
OVARC1000649	9.69	14.8	53.54	53.32	64.51	52.67		
OVARC1000661	1.14	9.33	2.99	5.34	5.24	5.24		
OVARC1000677	8.53	10.16	14.87	11.77	10.98	15.47		
OVARC1000678	1.21	4.49	2.71	3.28	4.17	3.26		
OVARC1000679	2.86	3.25	4.09	5.29	5.16	6.25	*	+
OVARC1000681	1.47	1.55	3.2	2.41	2.22	1.71		
OVARC1000682	10.09	11.33	50.91	33.79	47.49	44.31		
OVARC1000689	3.81	7.1	19.94	20.18	22.12	21.83		
OVARC1000700	1.8	10.37	3.18	4.98	4.37	5.14		
OVARC1000703	1.74	7.18	5.35	6.4	6.64	7.77		
OVARC1000722	10.59	11.92	47.93	43.41	60.06	39.34		
OVARC1000726	1.44	3.48	4.62	4.88	5.89	3.58		
OVARC1000727	1.93	2.09	4.13	3.78	3.79	3.89		
OVARC1000730	5.95	5.86	9.01	4.07	4.16	5.62		
OVARC1000741	4.85	6.13	8.74	15.19	10.58	13.71	*	+
OVARC1000746	0.89	3.61	2.43	2.06	2.9	2.84		
OVARC1000764	1.76	4.93	4.77	5.35	7.01	5.44		
OVARC1000769	1.13	4.3	3.6	3.76	4.42	5.2		
OVARC1000771	2.42	6.28	2.3	4.02	4.81	3.71		
OVARC1000773	19.09	24.7	31.93	44.69	56.24	46.24	*	+
OVARC1000775	11.67	8.94	16.44	12.16	8.7	4.26		
OVARC1000778	2.37	3.89	5.69	4.59	6.23	4.92		
OVARC1000779	0.8	2.02	1.85	2.23	2.45	1.46		
OVARC1000781	1.67	5.05	4.16	6.37	3.45	5.07		
OVARC1000787	1.64	4.79	4.22	2.97	5.44	3.25		
OVARC1000789	7.62	14.23	16.39	24.95	29.69	25.94	**	+
OVARC1000800	2.91	10.72	5.72	6.41	10.65	6.2		
OVARC1000802	1.55	8.77	2.97	2.99	5.34	2.76		
OVARC1000810	3.37	3.54	8.29	6.66	7.99	7.21		
OVARC1000811	2.41	2.73	7.5	4.88	3.67	4.95		
OVARC1000814	3.44	4.55	9.03	7.92	11.3	9.05		
OVARC1000816	7.64	10.41	12.41	10.99	10.58	14.11		
OVARC1000817	1.18	3.38	1.27	1.71	2.14	1.55		
OVARC1000834	2.46	8.3	3.39	4.84	5.81	4.01		
OVARC1000846	2.23	10.02	5.35	7.38	9.66	7.72		
OVARC1000850	1.74	8.37	3.38	3.39	2.51	3.1		
OVARC1000853	23.21	24.23	43.4	30.67	37.81	18.39		
OVARC1000862	2.28	2.66	4.91	3.05	1.61	1.49		
OVARC1000873	2.56	2.98	4.14	4.4	3.79	3.85		
OVARC1000875	1.47	3.07	1.79	2.35	3.09	1.87		
OVARC1000876	3.71	5.67	4.46	5.11	6.06	5.45		
OVARC1000883	6.06	9.53	9.18	12.43	15.42	13.05	*	+
OVARC1000885	2.84	9.95	3.38	3.74	5.66	4.11		
OVARC1000886	4.31	8.19	4.29	3.74	5.15	4.39		

(1160)

特開 2002-191363

2317

2318

OVARC1000890	17.47	18.3	91.22	70.97	78.71	51.68		
OVARC1000891	1.28	1.44	3.03	2.85	2.19	3.22		
OVARC1000897	0.48	1.74	1.29	1.21	1.33	0.5		
OVARC1000912	2.06	3.22	4.33	5.21	6.1	5.86	*	+
OVARC1000914	1.48	6.18	1.61	3.68	3.02	2.18		
OVARC1000915	1.71	6.64	4.14	4.87	3.54	4.76		
OVARC1000916	1.91	5.19	2.56	4.05	4.32	3.88		
OVARC1000924	1.45	5.5	3.09	3.28	3.85	3.48		
OVARC1000928	6.05	5.46	8.78	4.22	5.83	5.35		
OVARC1000936	1.37	1.39	2.17	2.04	3.25	2.49		
OVARC1000937	1.69	3.01	1.94	4.17	3.26	3.24		
OVARC1000945	1.55	3.67	3.62	3.83	3.22	2.28		
OVARC1000948	1.57	3.85	2.66	3.15	3.68	1.99		
OVARC1000956	1.41	5.08	4.36	7.56	7.26	5.51		
OVARC1000959	1.8	4.87	3.39	4.88	3.02	3.9		
OVARC1000960	2.64	7.53	9.55	11.64	13.89	12.86	*	+
OVARC1000964	19.89	17.19	103.98	118.41	165.46	96.14		
OVARC1000971	0.42	1.58	1.4	2.53	2.27	2.28	*	+
OVARC1000975	5.93	8.3	36.1	31.27	51.54	30.22		
OVARC1000976	0.65	2.12	1.27	2.17	1.46	1.5		
OVARC1000981	4.06	7.18	4.94	7.97	12.1	8.53		
OVARC1000982	2.83	5.41	2.23	3.13	3.02	3.54		
OVARC1000984	1.78	5.43	3.32	3.01	3.08	2.16		
OVARC1000995	2.94	6.59	4.5	5.98	6.19	6.72		
OVARC1000996	1.68	1.87	4.29	3.58	4.15	4.56		
OVARC1000999	6.02	5.65	15.29	15.61	13.18	13.29		
OVARC1001000	1.96	4.5	6.2	6.26	7.09	6.86		
OVARC1001004	0.51	3.4	1.45	2.05	3.3	1.47		
OVARC1001010	1.35	3.99	1.66	3.04	1.4	1.54		
OVARC1001011	1.46	5.57	1.13	2.39	3.27	2.45		
OVARC1001030	96.19	101.41	143.98	119.24	154.26	133		
OVARC1001032	1.42	5.34	1.89	1.83	2.82	1.92		
OVARC1001034	4.44	5.58	6.51	3.29	5.21	3.77		
OVARC1001038	3.62	5.03	7.4	10.3	10.88	8.61	*	+
OVARC1001040	2.63	3.77	6.93	5.25	6.51	4.25		
OVARC1001041	4.54	8.03	12.87	8.57	12.25	9.4		
OVARC1001044	1.05	2.92	1.83	1.96	2.43	1.84		
OVARC1001049	3.78	8.78	10.67	10.65	11.87	10.26		
OVARC1001051	40.95	55.97	80.66	66.89	109.71	87.49		
OVARC1001054	1.22	4.06	3.22	2.86	4.19	1.93		
OVARC1001055	2.13	3.38	3.82	4.32	5.61	5.22	*	+
OVARC1001062	5.8	6.15	12.54	8.04	9.94	9.57		
OVARC1001065	8.85	13.63	51.33	51.41	60.3	56.97		
OVARC1001068	2.82	5.62	4.76	4.72	4.02	5.52		
OVARC1001072	0.73	4.18	4.41	3.2	3.71	3.07		
OVARC1001073	0.92	5.7	2.65	2.91	2.54	1.79		
OVARC1001074	0.81	4.66	3.31	1.87	2.95	2.04		
OVARC1001078	2	5.12	2.79	3.57	3.08	2.83		
OVARC1001085	2.41	2.83	3.66	5.54	5.02	6.36	**	+
OVARC1001086	1.97	3.17	2.85	3.98	2.83	4.13		

(1161)

特開2002-191363

2319

2320

OVARC1001091	16.24	19.32	92.73	76.48	96.74	77.99		
OVARC1001092	4.62	5.35	7.22	9.69	7.84	6.05		
OVARC1001104	1.05	4.37	2.66	3.16	2.58	2.03		
OVARC1001107	11.59	15.6	40.28	31.21	49.49	42.22		
OVARC1001113	1.04	5.81	1.59	2.46	3.05	2.39		
OVARC1001117	2.71	6.63	4.31	4.67	5.74	2.67		
OVARC1001118	2.38	3.69	7.08	7.36	6.91	8.28		
OVARC1001125	2.02	2.9	3	5.92	4.97	5.9	**	+
OVARC1001129	2.61	4.58	3.19	8.46	9.43	11	**	+
OVARC1001132	1.7	6.48	2.66	3.69	4.26	4.66		
OVARC1001138	9.52	15.82	23.8	48.95	45.16	44.97	**	+
OVARC1001141	1.68	4.97	3.48	3.77	3.68	3.84		
OVARC1001154	18.31	29.49	68.39	60.43	83.49	65.64		
OVARC1001161	2.49	5.55	6.15	7.03	6.69	5.89		
OVARC1001162	2.2	3.13	5.34	5.09	4.86	5.26		
OVARC1001163	0.69	3.59	2.77	2.2	3.98	2.54		
OVARC1001167	3.03	4.57	7.69	10.19	12.95	9.3	*	+
OVARC1001169	0.74	4.87	2.68	2.47	1.91	2.06		
OVARC1001170	7.81	15.04	17.59	14.61	19.45	14.99		
OVARC1001171	15.57	17.71	24.31	16.12	23.34	22.51		
OVARC1001173	2.09	5.08	5.1	4.32	6.75	5.49		
OVARC1001176	22.57	21.48	89.96	76.74	102.3	70.7		
OVARC1001180	3.01	4.58	12.7	11.81	10.77	9.56		
OVARC1001188	2.66	3.7	3.95	3.62	3.44	4.16		
OVARC1001200	1.52	4.56	3.62	3.47	2.9	2.96		
OVARC1001202	3.75	6.65	6.53	9.26	7.79	10.23	*	+
OVARC1001206	1.52	5.52	1.15	1.59	1.13	1.9		
OVARC1001209	4.89	8.92	27.46	24.69	30.38	24.27		
OVARC1001219	1.81	6.36	4.71	5.33	3.95	3.62		
OVARC1001222	2.5	8.36	5.01	3.2	4.34	5.63		
OVARC1001232	2.91	4.18	7.74	6.75	6.02	5.65		
OVARC1001240	2.05	3.27	6.84	5.55	5.06	5.4		
OVARC1001243	0.94	2.59	1.76	3.64	2.64	1.86		
OVARC1001244	9.07	12.05	18	21.61	18.57	26.62		
OVARC1001246	30.48	50.95	48.51	80.54	100.83	101.88	**	+
OVARC1001247	3.64	9.86	7.7	6.57	7.02	4.49		
OVARC1001260	1.05	9.07	1.85	2.62	2.65	1.85		
OVARC1001261	4.23	10.5	6.99	3.46	2.08	2.94		
OVARC1001268	24.4	19.69	52.37	32.58	35.32	14.16		
OVARC1001270	14.46	15.1	20.83	9.69	9.8	8.65	*	-
OVARC1001271	2.62	3.62	3.88	3.95	7.02	4.26		
OVARC1001282	0.88	3.02	3.09	1.37	1.59	2		
OVARC1001296	3.02	8.06	2.3	3.04	4.11	5.41		
OVARC1001306	1.48	8.27	2.4	2.04	2.29	3.82		
OVARC1001314	0.49	8.47	1.57	1.06	1.79	1.32		
OVARC1001316	2.77	7.17	4.81	5.48	8.11	5.36		
OVARC1001329	6.12	6.18	21.11	17.09	19.29	16.22		
OVARC1001330	0.2	1.89	1.38	1.22	1.35	1.42		
OVARC1001336	1.92	3.7	3.59	5.67	4.09	4.02		
OVARC1001338	0.26	2.87	0.86	2.49	1.71	1.07		

(1162)

特開2002-191363

2321

2322

OVARC1001339	12.07	18.29	22.73	33.65	32.72	37.29	**	+
OVARC1001340	0.72	4.83	1.23	1.33	2.44	1.3		
OVARC1001341	4.35	9.25	6.77	7.94	11.38	9.69		
OVARC1001342	90.37	98.53	136.12	129.68	163.22	127.78		
OVARC1001344	2.1	2.51	6.27	6.52	6.89	6.2		
OVARC1001357	5.61	8.93	16.02	15.52	11.34	11.69		
OVARC1001359	8.96	12.4	16.15	21.66	13.84	10.6		
OVARC1001360	0.44	2.52	0.99	1.97	2.6	1.62		
OVARC1001369	1.56	5.66	1.89	3.41	1.88	3.1		
OVARC1001372	0.96	4.23	3.33	1.52	2.77	1.95		
OVARC1001376	1.82	5.1	3.62	5.79	5.79	4.18		
OVARC1001381	4.51	6.44	9.94	10.95	12.91	11.21		
OVARC1001391	0.5	1.62	1.44	1.88	1.27	1.26		
OVARC1001392	2.12	4.69	6.14	11.96	12.7	7.79	*	+
OVARC1001399	0.98	3.59	2.16	1.77	2.54	1.1		
OVARC1001417	1.01	3.07	1.76	2.39	3.61	2.81		
OVARC1001419	2.47	5.4	3.06	4.39	4.03	3.16		
OVARC1001425	2.29	5.58	5.15	8.76	8.5	8.07	*	+
OVARC1001436	1.37	5.85	2.54	2.9	3.57	3.18		
OVARC1001442	0.64	4.84	1.39	2.27	1.52	0.69		
OVARC1001451	3.09	2	3.89	5.18	5.98	4.75	*	+
OVARC1001452	1.35	2.41	2.87	2.96	4.69	3.13		
OVARC1001453	1.21	2.84	1.88	2.3	1.82	1.57		
OVARC1001476	10.67	14.38	16.52	17.22	12.9	13.09		
OVARC1001480	0.93	4.73	1.5	3.1	2.98	2.21		
OVARC1001489	0.97	6.89	2.51	3.01	2.83	2.09		
OVARC1001493	2.09	6.59	3.75	7.38	8.78	10.48	*	+
OVARC1001496	4.65	9.58	8.63	10.74	7.37	11.03		
OVARC1001499	1.24	1.18	2.6	3.47	2.68	2.84		
OVARC1001506	2.9	2.7	5.31	5.33	6.73	5.48		
OVARC1001509	1.73	3.89	3.69	4.07	4.59	3.07		
OVARC1001510	0.16	3.69	1.42	1.94	1.73	0.86		
OVARC1001516	2.57	5.78	3.85	6.04	4.97	4.39		
OVARC1001525	0.53	4.76	2.12	1.94	2.01	1.81		
OVARC1001542	8.8	12.86	13.01	15.91	13.63	17.23		
OVARC1001544	2.14	6.6	6.72	7.54	8.33	6.22		
OVARC1001546	4.08	4.32	4.6	6.12	5.31	7.23	*	+
OVARC1001547	1.29	2.53	1.68	2.44	1.85	2.22		
OVARC1001555	10.39	16.51	68.77	48.66	65.39	56.39		
OVARC1001560	3.35	4.91	5.52	5.36	4.93	5.34		
OVARC1001569	1.63	4.75	4.79	5.92	5.19	5.1		
OVARC1001570	3.96	7.9	6.93	7.72	10.7	8.55		
OVARC1001577	1.68	5.89	5.41	8.61	6.9	10.2		
OVARC1001578	0.25	3.47	-0.19	-0.47	0.24	0.15		
OVARC1001596	12.13	11.65	14.23	13.51	14.82	27.15		
OVARC1001600	1.13	2.9	1.48	2.81	2.67	3.67		
OVARC1001607	6.22	7.72	10.91	13.42	14.01	13.45	*	+
OVARC1001610	1.81	5.25	2.84	4.25	2.66	2.7		
OVARC1001611	0.13	5.11	1.24	1.48	2.89	1.79		
OVARC1001615	0.58	5.42	1.93	1.54	2.56	1.74		

(1163)

特開 2002-191363

2323

2324

OVARC1001636	1.09	3.75	1.05	2.39	2.05	2.15		
OVARC1001668	3.77	6.75	10.04	10.5	11.4	10.48		
OVARC1001702	1.18	2.21	2.42	3.86	2.07	2.25		
OVARC1001703	2.82	3.18	2.97	2.64	4.71	4.65		
OVARC1001710	3.58	7.03	8.67	8.01	6.28	9.55		
OVARC1001711	1.96	7.3	3.36	5.01	4.15	5.4		
OVARC1001713	9.17	11.54	44.65	36.47	51.48	43.79		
OVARC1001725	1.01	5.45	6.11	2.56	3.77	4.01		
OVARC1001726	1.64	4.48	3.23	4.97	5.6	5.18		
OVARC1001727	1.4	2.41	1.52	1.43	2.35	1.14		
OVARC1001731	120.62	110.86	255.43	140.73	139.03	74.2		
OVARC1001735	1.29	3.44	3.54	3.75	3.25	2.89		
OVARC1001741	3.3	4.73	15.28	13.09	12.93	13.17		
OVARC1001745	2.72	5.39	6.83	9.17	10.23	8.89	*	+
OVARC1001759	3.31	9.01	6.31	7.61	7.61	11.95		
OVARC1001762	3.96	7.78	6.38	10.3	11.01	13.4	*	+
OVARC1001766	5.33	7.8	11.99	15.56	16.86	15.33	*	+
OVARC1001767	0.94	3.76	1.18	1.97	1.96	2.41		
OVARC1001768	3.31	3.86	3.76	5.35	4.59	3.54		
OVARC1001770	3.04	6.58	9.98	11.46	9.08	9.46		
OVARC1001776	2.11	4.7	3.5	4.72	3.64	2.92		
OVARC1001791	1.13	4.77	3.54	3.07	3.12	3.42		
OVARC1001795	0.89	6.19	1.24	2.31	2.87	2.11		
OVARC1001798	2.81	12.11	7.57	9.72	11.93	9.04		
OVARC1001802	1.73	11.64	4.9	5.6	5.93	4.01		
OVARC1001805	1.92	6.96	2.58	3.62	4.59	3.51		
OVARC1001807	1.9	2.53	4.18	3.06	3.12	2.46		
OVARC1001809	12.38	14.06	76.32	55.87	81.41	52.83		
OVARC1001812	1.44	3.39	3.15	3.23	4.63	3.71		
OVARC1001813	1.61	4.29	2.33	2.93	3.98	2.51		
OVARC1001820	1.67	7.15	3.21	3.47	3.76	3.22		
OVARC1001828	0.78	6.85	2.36	1.91	3.23	2.2		
OVARC1001833	1.07	8.12	2.02	2.4	2.1	1.92		
OVARC1001839	1.56	8.43	2.98	2.91	3.59	1.15		
OVARC1001846	1.91	1.38	2.9	2.15	2.11	1.8		
OVARC1001849	1.21	2.52	2.42	5.79	3.69	4.03	*	+
OVARC1001861	1.46	3.56	2.73	2.78	2.5	2.09		
OVARC1001873	3.09	3.78	4.68	5.47	4.42	4.73		
OVARC1001879	1.44	6.08	3.48	2.35	2.93	2.22		
OVARC1001880	0.91	7.84	2.94	3.63	5.78	3.8		
OVARC1001883	0.99	7.61	3.12	2.61	3.42	2.52		
OVARC1001900	1.11	7.07	4.03	2.57	3.61	2.89		
OVARC1001901	0.54	1.84	1.21	2.42	1.37	2.43		
OVARC1001911	0.59	1.57	1.66	1.51	1.39	1.55		
OVARC1001916	1.86	3.13	3.35	4.16	4.61	3.51		
OVARC1001928	1.45	3.53	1.55	1.58	1.62	1.72		
OVARC1001937	5.12	11.69	8.13	17.41	11.63	15.16		
OVARC1001940	1.1	4.51	3.13	3.72	3.14	2.78		
OVARC1001942	3.85	7.4	8.03	11.47	13.91	12.77	*	+
OVARC1001943	7.16	10.07	11.08	9.62	13.85	11.87		

(1164)

特開 2002-191363

2325

2326

OVARC1001949	1.69	3.34	4.15	5.35	4.01	5.55		
OVARC1001950	1.53	2.41	3.79	6.3	4.35	3.98		
OVARC1001952	11.3	11.38	53.57	52.33	78.84	38.05		
OVARC1001954	1.12	2.99	2.2	3.09	2.67	2.05		
OVARC1001963	1	4.91	2.89	4.5	3.39	3.21		
OVARC1001983	3.62	14.16	14.25	20.96	19.21	21.67		
OVARC1001987	3.12	6.54	5.94	6.08	8.39	8.02		
OVARC1001989	1.41	5.2	4.96	4.54	5.59	5.26		
OVARC1001991	1.74	3.27	4.08	4.57	3.86	3.27		
OVARC1002005	4.14	3.55	7.66	10.01	9.06	8.2	*	+
OVARC1002044	3.73	3.94	6.17	6.57	8.32	6.99	*	+
OVARC1002046	10.28	16.21	20.07	29.4	37.78	37.02	**	+
OVARC1002050	1.7	5.6	2.43	3.96	3.82	2.53		
OVARC1002058	4.23	6.11	4.02	4.69	5.55	5.43		
OVARC1002066	11.47	13.5	25.49	26.02	28.69	22.63		
OVARC1002082	3.6	8.55	8.81	9.6	8.89	6.49		
OVARC1002091	3.17	5.67	8.37	9.49	5.49	7.64		
OVARC1002092	1.38	2.72	2.2	4	3.97	1.88		
OVARC1002093	1.79	3.1	4.51	5.01	4.44	3.88		
OVARC1002094	1.55	6.24	4.17	36.42	27.25	28.35	**	+
OVARC1002107	1.42	4.63	2.69	4.86	5.48	3.6		
OVARC1002112	6.17	11.59	8.5	13.47	17.48	11.92		
OVARC1002126	2.66	6.35	6.68	7.95	6.44	8.79		
OVARC1002127	0.73	5.04	1.86	1.92	2.61	1.52		
OVARC1002138	1.4	1.79	1.86	3.16	4.82	2.75	*	+
OVARC1002143	0.73	1.51	1.55	1.29	3.03	2.09		
OVARC1002156	2.42	3.87	4.19	4.43	3.9	3.65		
OVARC1002158	0.88	2.63	1.6	2.36	1.57	1.51		
OVARC1002165	4.85	6.3	9.83	10.73	14.03	10.87		
OVARC1002176	0.86	5.08	3.59	2.01	3.46	2.64		
OVARC1002178	0.83	5.35	3.12	3.8	5.02	4.25		
OVARC1002182	1.29	2.89	3.77	2.45	4.64	3.12		
OVARC1002185	11.45	13.19	62.79	43.91	53.43	55.56		
PLACE1000004	1.42	3.23	2.35	3.87	4.25	4.05	*	+
PLACE1000005	1.18	3.06	3.3	5.27	5.31	4.83	*	+
PLACE1000006	2.01	8.33	3.23	4.2	5.44	4.67		
PLACE1000007	0.97	5.13	2.89	3.03	2.47	2.56		
PLACE1000014	2.9	8.06	6.26	6.67	8.18	6.55		
PLACE1000031	0.88	4.81	0.45	2.61	2.71	2.79		
PLACE1000033	1.23	2.15	2.75	2.42	3.17	2.56		
PLACE1000040	3.08	4.43	6.18	7.11	5.54	7.37		
PLACE1000048	1.83	3.24	2.14	3.32	3.96	3.74		
PLACE1000050	2.12	5.36	9.1	9	6.55	8.25		
PLACE1000061	138.29	147.36	249.77	165.55	233.98	230.37		
PLACE1000066	14.23	15	19.46	15.86	15.62	18.52		
PLACE1000075	3.03	6.24	9.08	4.98	6.93	7.11		
PLACE1000078	2.1	5.75	5	6.07	6.93	5.19		
PLACE1000081	1.08	1.88	1.52	1.13	1.89	1.27		
PLACE1000086	4.97	6.55	11.25	8.1	9.16	7.75		
PLACE1000094	0.7	4.18	1.72	1	3.44	2.96		

(1165)

特開2002-191363

2327

2328

PLACE1000101	4.67	8.44	7.7	11.69	10.38	13.65	*	+
PLACE1000121	0.87	6.29	2.02	1.95	2.85	2.39		
PLACE1000133	6.65	11.93	17.66	15.19	17.59	21.71		
PLACE1000142	1.79	6.03	5.66	2.64	4.77	4.24		
PLACE1000146	1.95	4.51	2.89	3.71	5.02	2.82		
PLACE1000163	4.52	5.99	10.71	16.27	10.95	13.78		
PLACE1000172	1.12	2.63	1.48	1.81	3.21	2.29		
PLACE1000181	1.06	3	2.98	2.63	3.75	2.86		
PLACE1000184	1.17	3.08	1.48	1.87	3.28	2.27		
PLACE1000185	2.99	6.52	8.47	9.53	9.99	12.03		
PLACE1000198	0.78	4.09	2.49	2.19	2.17	1.93		
PLACE1000213	3.3	5.87	7.36	4.35	5.38	8.09		
PLACE1000214	1.37	4.29	4.54	5.22	6.72	4.22		
PLACE1000220	9.61	7.84	16.78	7.48	5.77	4.1		
PLACE1000231	2.48	4.1	4.92	4.57	4.65	3.91		
PLACE1000236	0.66	3.33	2.43	2.8	3.63	2.37		
PLACE1000245	2.92	5.88	6.34	9.34	11.24	10.55	*	+
PLACE1000246	5.15	8.27	9.95	3.29	2.55	2.86	*	-
PLACE1000258	5.4	12.61	13.52	14.88	16.7	14.95		
PLACE1000288	1.68	7.22	2.96	2.83	4.02	2.48		
PLACE1000292	3.72	9.02	8.85	10.23	20.58	9.21		
PLACE1000302	0.56	1.01	1.39	1.07	0.92	0.62		
PLACE1000304	1.13	3.26	3.17	3.75	2.32	3.05		
PLACE1000308	2.54	4.35	4.17	4.42	3.87	1.34		
PLACE1000309	2.29	4.02	4	6.72	5.23	7.88	*	+
PLACE1000312	1.33	3.44	2.48	2.74	3.99	2.5		
PLACE1000330	0.46	5.76	3.02	1.32	1.93	1.35		
PLACE1000332	1.02	8.82	2.01	3.01	3.78	1.68		
PLACE1000347	2.3	9.48	3.89	2.59	5.81	3.22		
PLACE1000351	1.2	1.5	2.87	2.2	2.4	2		
PLACE1000374	2.01	3.03	7.02	8.89	6.55	6.85		
PLACE1000380	2.39	4.27	3.95	4.9	2.12	2.38		
PLACE1000383	1.03	2.62	1.9	2.53	3.64	2.4		
PLACE1000397	0.63	4.06	1.89	2.82	3.34	3.47		
PLACE1000401	1.22	6.39	2.24	2.23	3.05	2.36		
PLACE1000406	1.08	8.76	3.4	3.72	4.08	3.64		
PLACE1000412	1.61	6.38	1.56	1.62	3.45	1.46		
PLACE1000420	2.59	3.51	4.6	8.95	7.28	5.6	*	+
PLACE1000421	0.99	1.3	2.32	2.97	2	1.7		
PLACE1000423	16.6	23.29	32.85	10.67	8.02	8.09	*	-
PLACE1000424	1.36	3.09	2.12	3.35	2.65	1.97		
PLACE1000430	0.77	4.36	1.95	3.51	2.94	3.77		
PLACE1000433	1.06	5.9	1.65	1.89	2.6	1.8		
PLACE1000435	1.39	7.21	4.77	6.22	6.29	4.22		
PLACE1000437	6.06	10.65	10.14	17.29	20.07	18.79	**	+
PLACE1000442	3.75	3.85	6.27	7.81	5.7	6.7		
PLACE1000444	2.14	3.94	8.96	11.14	11.55	9.8		
PLACE1000453	5.57	11.03	14.16	10.42	7.4	2.99		
PLACE1000456	1.25	2.21	1.97	1.33	2.18	1.07		
PLACE1000465	2.09	5.63	6.62	12.97	11.8	10.69	**	+

(1166)

特開 2002-191363

2329

2330

PLACE1000481	2.32	8.1	3.73	6.89	6.64	6.45		
PLACE1000492	1.15	4.45	2.95	3.27	3.06	2.81		
PLACE1000508	1.36	4.64	4	3.91	4.24	3.71		
PLACE1000512	4.91	4.29	6.39	8.12	7.8	4.6		
PLACE1000540	5.18	3.93	7.84	5.44	6.9	5.57		
PLACE1000541	13.59	15.07	48.84	60.62	81.24	41.96		
PLACE1000546	0.86	3.61	2.82	4.72	4.63	2.5		
PLACE1000547	2.16	4.61	3.83	6.31	5.64	5.92	*	+
PLACE1000560	2.08	5.97	2.1	1.62	2.8	1.72		
PLACE1000562	2.8	6.23	6.04	8.86	11.26	8.61	*	+
PLACE1000564	1.54	6.4	3.07	3.16	4.41	3.43		
PLACE1000583	3.75	3.28	6.32	6.78	11.53	6.8		
PLACE1000587	8.52	9.32	12.99	13.64	14.69	9.43		
PLACE1000588	1.92	4.36	3.99	8.79	8.15	4.48		
PLACE1000596	1.99	5.34	4.39	7.8	6.74	4.51		
PLACE1000599	2.39	5.51	7.05	7.92	7.79	6.46		
PLACE1000605	5.12	11.43	7.06	14.2	15.1	12.67	*	+
PLACE1000610	2.01	6.08	3.54	5.26	4.48	2.94		
PLACE1000611	13.18	19.13	24.68	16.45	20.37	22.79		
PLACE1000626	3.19	4.04	8.04	5.71	9.93	8.12		
PLACE1000633	1.32	2.12	3.95	4.8	6.06	3.59		
PLACE1000636	1.15	2.54	1.43	2.72	4.25	2.94		
PLACE1000653	5.07	8.56	9.29	11.07	11.87	14.08	*	+
PLACE1000656	4.2	12.9	25.22	16.66	16.71	12.92		
PLACE1000663	2	6.43	3.59	2.39	6.61	4.03		
PLACE1000706	2	6.3	5.04	5.37	7.27	6.26		
PLACE1000712	3.9	9.52	10.82	10.49	10.07	9.11		
PLACE1000716	0.98	2.75	3.44	2.44	3.2	2.82		
PLACE1000740	2.74	5.28	6.24	8.83	8.13	9.69	*	+
PLACE1000748	3.35	3.51	6.81	3.12	5.02	4.23		
PLACE1000749	3.49	6.35	5.94	4.61	4.65	6.02		
PLACE1000751	2.71	5.34	4.07	7.81	8.32	8.36	**	+
PLACE1000755	1.39	6.14	1.93	2.55	5.1	2.96		
PLACE1000769	2.29	6.8	3.45	3.33	4.58	2.6		
PLACE1000778	0.87	1.48	1.99	2.05	2.94	2.38		
PLACE1000785	9.56	12.21	27.18	28	24.34	29.54		
PLACE1000786	2.68	4.22	3.63	3.09	3.77	3.7		
PLACE1000793	4.05	7.21	6.7	6.06	7.6	9.1		
PLACE1000795	2.15	5.5	3.99	4.44	5.29	4.31		
PLACE1000798	0.88	8.44	3.24	3.13	3.8	3.72		
PLACE1000812	2.13	5.08	4.46	5.06	5.16	6.03		
PLACE1000823	1.71	5.2	4.89	5.67	7.28	4.84		
PLACE1000825	1.6	2.86	2.02	3.77	3.96	3.76	*	+
PLACE1000838	16	15.77	23.73	13.88	15.6	15.65		
PLACE1000841	1.22	3.78	3.31	3.97	10.65	3.77		
PLACE1000843	2.14	6.2	5.68	5.79	7.7	5.38		
PLACE1000849	2.79	8.82	6.72	7.24	6.78	10.02		
PLACE1000856	2.01	5.3	3.59	3.42	4.79	4.19		
PLACE1000863	5.2	7.58	9.56	8.97	12.34	11.53		
PLACE1000876	3.65	7.6	6.02	6.7	9.95	9.06		

(1167)

特開2002-191363

2331

2332

PLACE1000899	1.36	2.24	3.12	4.12	5.14	4.22	*	+
PLACE1000907	4.82	5.53	9.59	6.77	8.44	5.83		
PLACE1000909	1.18	3.31	2.45	3.65	3.88	3.44		
PLACE1000912	0.42	4.55	1.77	1.76	2.72	1.46		
PLACE1000914	1.05	4.41	3.5	3	6.09	4.22		
PLACE1000918	0.54	4.49	1.61	1.82	3.13	1.98		
PLACE1000927	10.48	12.41	16.9	20.91	23.21	25.47	*	+
PLACE1000931	0.69	3.44	2.12	2.44	3.94	3.3		
PLACE1000944	2.55	2.24	4.78	3.84	3.32	2.09		
PLACE1000948	0.52	2.31	2.96	2.21	2.72	1.72		
PLACE1000958	0.12	2.2	1.73	1.11	1.77	2.27		
PLACE1000972	1.01	3.43	2.89	4.49	5.33	3.75		
PLACE1000977	2.33	5.67	4.42	2.71	5.33	5.25		
PLACE1000979	1.63	8.01	3.93	4.24	5.92	4.57		
PLACE1000986	3.37	16.51	6.63	6.97	8.75	7.69		
PLACE1000987	1.76	10.13	4.79	4.17	4.74	5.11		
PLACE1001000	4.85	4.62	7.76	6.02	4.25	3.02		
PLACE1001007	7	6.94	14.66	5.39	3.76	3.47		
PLACE1001010	0.61	2.04	2.45	2.56	2.73	2.84		
PLACE1001015	0.88	2.55	1.84	2.36	1.72	2.42		
PLACE1001016	1.79	4.54	4.29	6.37	9	6.57	*	+
PLACE1001022	0.68	6.5	2.45	1.9	2.39	1.29		
PLACE1001024	1.05	8.89	1.83	1.34	2.49	2.35		
PLACE1001036	2.63	10.55	5.42	3.62	5.49	5.43		
PLACE1001038	50.16	49.81	118.83	82.67	64.83	52.8		
PLACE1001048	1.07	1.82	0.92	2.39	2.09	1.21		
PLACE1001054	9.95	10.74	63.88	62.96	79.44	66.71		
PLACE1001062	1.45	4.19	3.71	3.12	4.17	3.05		
PLACE1001063	1.35	3.74	2.2	4.06	3.65	2.74		
PLACE1001076	0.46	6.48	1	1.86	2.18	1.62		
PLACE1001081	1.53	7.95	3.33	3.65	5.24	4.8		
PLACE1001088	1.32	5.24	1.22	1.42	2.81	1.37		
PLACE1001092	2.31	2.47	4.68	5.8	4.18	3.9		
PLACE1001098	0.93	2.62	4.53	5.49	4.05	3.17		
PLACE1001100	1.31	2.58	2.48	4.27	4.17	3.37	*	+
PLACE1001104	1.95	4.09	4.54	3.39	3.42	4.47		
PLACE1001114	1.56	6.54	4.33	5.17	3.78	3.27		
PLACE1001118	2.52	5.77	6.12	6.21	6.14	5.1		
PLACE1001123	2.86	5.3	7.53	7.08	8.51	7.63		
PLACE1001136	1.58	4.39	5.13	5.29	5.95	5.85		
PLACE1001144	6.27	5.67	13.43	10.34	11	10.08		
PLACE1001147	2.11	2.98	6.03	6.13	5.15	4.83		
PLACE1001148	1.72	1.31	1.89	1.76	2.39	2.31		
PLACE1001159	0.86	2.37	2.05	2.27	3.73	1.43		
PLACE1001168	8.87	14.52	15.09	25.46	23.18	30.79	*	+
PLACE1001171	0.69	3.89	1.23	2.53	1.42	1.53		
PLACE1001183	0.24	3.61	1.81	1.57	2.78	1.38		
PLACE1001185	3.13	7.43	3.76	5	6.4	5.64		
PLACE1001201	1.77	2.8	3.29	6.32	6.94	6.32	**	+
PLACE1001229	7.51	8.56	12.64	15.24	11.45	10.42		

(1168)

特開 2002-191363

2333

2334

PLACE1001231	1.83	2.73	3.07	4.09	5.1	2.3		
PLACE1001238	1.52	4.35	3.74	3.65	4.52	4.57		
PLACE1001241	1.63	5.58	2.92	5.73	8.13	7.04		
PLACE1001242	22.28	29.54	30.28	46.43	48.89	62.65	*	+
PLACE1001247	2.43	7.02	4.07	5.03	5.91	4.52		
PLACE1001250	1.01	5.36	3.61	4.68	4.39	4.81		
PLACE1001257	2.99	3.06	7.06	7.89	9.21	7.69		
PLACE1001272	3.19	4.27	5.68	7.13	6.43	5.14		
PLACE1001279	0.96	3.12	2.74	3.08	3.81	3.29		
PLACE1001280	1.08	4.75	2.68	4.98	4.45	2.86		
PLACE1001294	1.91	7.23	6.91	4.88	5.57	6.18		
PLACE1001295	4.16	9.94	7.53	8.55	11.85	8.43		
PLACE1001300	2.46	7.9	4.31	4.65	14.73	4.95		
PLACE1001304	3	8.27	10.47	8.57	10.81	10.64		
PLACE1001311	3.95	3.34	5.67	6.85	9.14	7.6	*	+
PLACE1001323	2.17	2.95	5.12	5.66	8.43	5.5		
PLACE1001325	0.88	1.95	3.71	2.84	3.56	3.27		
PLACE1001340	5.18	6.99	9.8	8.69	12.02	10.48		
PLACE1001344	1.52	3.49	1.77	2.34	2.06	1.75		
PLACE1001351	3.23	6.39	8.39	6.4	8.62	6.1		
PLACE1001366	1	4.49	4.02	4.19	4.6	3.72		
PLACE1001377	3.02	4.97	5.1	7.91	7.28	5.83	*	+
PLACE1001383	2.31	4.13	3.53	2.62	5.5	5.72		
PLACE1001384	1.81	3.23	2.89	2.05	3.43	3.15		
PLACE1001387	1.65	3.64	3.7	3.03	4.83	3		
PLACE1001395	3.72	6.64	6.54	7.01	7.61	7.73		
PLACE1001399	3.71	6.58	9.31	7.37	8.61	9.22		
PLACE1001401	0.83	5.25	2.33	1.55	1.76	0.87		
PLACE1001407	11.65	21.8	24.47	22.63	18.09	26.24		
PLACE1001412	1.6	4.98	4.53	4.08	4.42	3.83		
PLACE1001414	2.3	3.02	5.86	7.57	5.13	6.83		
PLACE1001416	2.99	4.71	3.29	5.62	4.04	7.08		
PLACE1001433	33.62	33.05	51.64	49.1	58.33	55.88		
PLACE1001440	1.95	3.99	3.96	3.6	3.53	2.1		
PLACE1001456	1.64	5.5	4.26	4.15	4.87	4.49		
PLACE1001464	32.76	28.05	47.41	53.22	68.42	61.32	*	+
PLACE1001468	0.85	5.04	1.17	1.56	2.55	2.27		
PLACE1001484	1.31	4.85	2.96	4.25	5.8	3.04		
PLACE1001500	0.92	2.22	2.14	2.72	3.34	3.26	*	+
PLACE1001502	1.36	3.6	3.9	3.54	5.9	4.54		
PLACE1001503	1.7	4.58	6.72	7.47	8.2	8.05		
PLACE1001505	6.34	14.13	16.16	39.97	27.14	46.65	*	+
PLACE1001513	4.09	10.82	8.17	5.87	8.53	14.61		
PLACE1001516	0.61	4.33	1.33	1.71	3.49	1.99		
PLACE1001517	5.56	8.58	14.77	14.14	14.96	14.28		
PLACE1001523	12.83	14.09	20.42	22.79	19.74	32.9		
PLACE1001526	5.12	4.89	8.42	9.51	9.11	6.89		
PLACE1001534	2.12	5.12	3.58	3.62	5.55	3.99		
PLACE1001536	0.61	2.5	1.52	2.11	3.2	1.9		
PLACE1001545	17.97	23.9	38.46	33.78	45.13	66.08		

(1169)

特開 2002-191363

2335

2336

PLACE1001551	2.55	6.26	6.15	4.72	6.59	6.71		
PLACE1001564	1.37	4.87	2.88	4.01	3.57	3.7		
PLACE1001570	2.62	5.95	4.18	2.19	3.82	4.32		
PLACE1001571	2.04	4.51	6.07	5.69	6.27	5.81		
PLACE1001595	4.73	4.64	10.04	11.6	8.27	5.28		
PLACE1001602	7.23	8.39	18.65	20.38	18.68	19.71		
PLACE1001603	2.01	3.83	5.37	6.86	5.86	4.56		
PLACE1001608	3.44	7.22	5.9	5.82	7.73	8.7		
PLACE1001610	3.77	8.4	8.22	9.26	9.49	9.85		
PLACE1001611	1.94	7.34	3.65	2.28	3.85	1.88		
PLACE1001629	0.78	6.77	2.24	3.62	3.36	3.52		
PLACE1001632	1.66	8.26	4.04	4.3	4.14	4.37		
PLACE1001634	7.4	9.92	39.12	23.85	32.41	18.38		
PLACE1001637	0.84	2.16	1.25	1.41	2.4	1.1		
PLACE1001640	1.33	3.27	4.66	2.68	4.85	4.49		
PLACE1001655	0.83	2.93	2.06	2.82	2.14	2.02		
PLACE1001672	1.84	7.04	4.01	3.3	4.41	4.09		
PLACE1001676	1.38	8.49	3.54	4.63	4.77	3.85		
PLACE1001683	12.79	23.62	24.61	25.33	30.22	27.13		
PLACE1001691	3.41	12.29	6.72	9.03	8.96	9.83		
PLACE1001692	1.47	2.96	5.25	5.87	5.6	5.13		
PLACE1001705	3.02	3.75	9.88	10.06	9.21	8.32		
PLACE1001716	1.68	3	2.61	2.24	3.79	3.58		
PLACE1001720	1.49	2.62	2.21	1.56	2.45	1.71		
PLACE1001728	1.43	6.19	4.24	1.96	2.04	2.51		
PLACE1001729	2.12	8.13	4.44	3.8	4.52	4.36		
PLACE1001739	2.61	9.55	4.04	4.95	7.24	6.16		
PLACE1001740	0.92	5.36	2.09	1.92	2.1	1.69		
PLACE1001745	1.15	0.98	3.22	1.87	2.48	2.31		
PLACE1001746	1.04	2.25	2.55	4.64	3.4	2.82		
PLACE1001748	4.74	7.01	8.18	8.19	6.58	5.96		
PLACE1001753	2.06	3.54	3.29	7.44	5.57	5.82	*	+
PLACE1001756	5.6	11.31	38.07	31.78	44.99	35.99		
PLACE1001760	6.54	12.23	12.85	16.36	16.96	16.66	*	+
PLACE1001767	11.26	14.98	59.72	45.37	61.46	45.39		
PLACE1001771	1.96	6.64	4.03	4.32	5.22	4.54		
PLACE1001775	2.23	2.81	6.72	5.1	3.11	4.79		
PLACE1001777	83.34	145.9	190.82	142.92	71.27	59.69		
PLACE1001781	1.9	3.86	4.91	8.72	3.39	2.3		
PLACE1001783	0.76	3.21	2.06	4.84	2.09	1.54		
PLACE1001786	1.77	6.61	2.72	3.7	3.32	2.6		
PLACE1001788	5.16	9.07	7.14	10.52	8.74	9.32		
PLACE1001795	1.92	4.4	4.82	5.42	4.61	5.42		
PLACE1001799	0.69	3.62	2.11	1.86	2.83	1.97		
PLACE1001810	0.89	1.52	1.76	2.73	3.91	1.73		
PLACE1001817	5.53	6.12	10.88	10.56	9.4	6.38		
PLACE1001821	4.68	6.07	7.11	8.37	9.92	4.99		
PLACE1001836	0.91	3.12	2.38	2.69	4.12	2.63		
PLACE1001844	1.55	5.1	3.48	4.42	4.36	4.09		
PLACE1001845	3.62	8.38	7.39	7.88	7.55	9.93		

(1170)

特開2002-191363

2337

2338

PLACE1001858	2.56	6.58	3.52	5.26	7.48	6.05		
PLACE1001869	3.13	7.15	4.85	6.09	6.46	5.66		
PLACE1001890	11.74	11.92	21.45	173.44	255.31	125.13	*	+
PLACE1001897	9.19	13.85	16.44	22.22	23.13	12.95		
PLACE1001902	10.13	12.6	21.53	22.74	27.67	12.77		
PLACE1001904	1.38	3.72	1.51	2.45	2.53	2.35		
PLACE1001907	3.36	6.76	5.71	7.67	5.67	5.59		
PLACE1001910	83.6	82.16	135.34	301.29	325.42	244.59	**	+
PLACE1001912	1.53	6.6	3.36	5.54	5.48	4.85		
PLACE1001918	17.31	22.95	30.16	31.14	40.44	40.02	*	+
PLACE1001920	2.07	3.51	5.43	11.97	13.8	11.4	**	+
PLACE1001928	3.06	2.96	4.67	5.29	9.7	5.31		
PLACE1001930	1.17	3.92	2.2	2.9	4.73	3.22		
PLACE1001949	1.16	3.67	1.78	3.84	4.24	3.18		
PLACE1001959	1.36	4.7	3.16	2.63	3.17	2.26		
PLACE1001969	2.09	7.83	7.21	6.56	10.73	6.57		
PLACE1001974	7.39	11.98	11.87	11.43	16.09	16.06		
PLACE1001981	0.77	4.38	3.22	1.77	3.88	2.36		
PLACE1001983	3.81	4.12	5.32	5.92	6.16	5.72	*	+
PLACE1001989	2.34	4.15	5.02	4.37	5.91	3.72		
PLACE1002004	3.07	4.06	8.05	9.22	9.69	7.18		
PLACE1002008	8.4	11.76	17	23.36	22.19	22.42	*	+
PLACE1002015	26.96	30.92	67.62	105.75	88.42	94.15	*	+
PLACE1002044	3.79	8.07	5.86	4.64	6.39	6.4		
PLACE1002046	1.78	5.68	1.9	4.3	5.79	4.97		
PLACE1002052	1.09	4.98	2.26	1.38	2.41	2.32		
PLACE1002066	4.79	6.3	8.29	10.24	10.77	9.93	*	+
PLACE1002072	2.55	3.91	4.86	6	5.48	6	*	+
PLACE1002073	0.51	2.83	2.29	2.35	4.06	2.91		
PLACE1002080	1.81	6.49	6.13	4.76	6.82	5.72		
PLACE1002081	1.66	6.13	4.06	3.74	4.86	4.3		
PLACE1002090	7.74	16.55	13.87	12.53	14.4	19.41		
PLACE1002095	2.97	6.22	8.45	10.01	10.18	11.32	*	+
PLACE1002102	4.26	8.56	8.81	9.47	9.56	10.67		
PLACE1002109	2.57	5.08	3.81	4.66	6.17	6.32		
PLACE1002115	1.75	3.57	2.48	2.78	4.26	2.59		
PLACE1002119	15.65	15.3	35.78	37.28	32.59	38.23		
PLACE1002140	5.25	8.45	14.05	19.93	17.14	17.19	*	+
PLACE1002150	1.54	8.26	4.25	3.23	5.36	4.12		
PLACE1002153	1.6	5.75	2.58	3.47	5.76	4.48		
PLACE1002157	0.87	2.96	1.72	1.76	3.28	3.73		
PLACE1002163	2.13	4.67	4.55	6.21	8.81	6.03		
PLACE1002168	2.39	4.04	4.42	4.68	6.28	3.22		
PLACE1002170	2.73	3.53	7.35	4.89	5.33	3.74		
PLACE1002171	3.09	6.11	10.02	18.44	14.63	13.93	*	+
PLACE1002180	3.16	6.23	6.77	4.63	8.54	8.39		
PLACE1002184	9.2	15.58	18.42	124.63	250.27	333.14	*	+
PLACE1002200	1.35	5.67	2.38	2.97	2.89	2.26		
PLACE1002205	3.3	6.47	18.07	17.08	18.38	14.61		
PLACE1002213	2.2	4.16	4.93	5.52	8.03	6.03		

(1171)

特開2002-191363

2339

2340

PLACE1002219	1.05	1.91	2.23	3.33	3.53	1.96		
PLACE1002227	0.68	2.65	1.7	1.5	3.03	1.67		
PLACE1002253	0.32	2.67	1.28	1.47	0.76	0.43		
PLACE1002256	1.16	4.78	3.31	3.54	3.01	4.79		
PLACE1002259	1.46	5.69	4.48	3.22	2.98	2.31		
PLACE1002285	1.16	10.74	2.29	1.55	2.38	1.24		
PLACE1002301	9.42	17.5	14.68	12.7	10.48	11.7		
PLACE1002310	4.28	10.16	9.86	8.82	7.87	9.94		
PLACE1002311	1.84	2.94	3.87	2.96	2.87	2.03		
PLACE1002319	2.31	2.64	2.94	3.21	3.23	3.92	*	+
PLACE1002329	0.56	2.54	2.5	4.07	3.58	3.07		
PLACE1002333	1.34	3.1	1.96	1.22	2.44	2		
PLACE1002342	4.19	9.04	9.44	5.06	8.52	8.17		
PLACE1002343	0.49	6.98	2.94	2.08	1.9	2.52		
PLACE1002355	1.31	9.39	2.36	3.33	4.35	2.63		
PLACE1002358	1.15	7.94	3.3	2.6	2.65	2.13		
PLACE1002359	1.91	2.17	3.47	4.7	3.91	3.42		
PLACE1002374	29.69	28.18	54.19	53.9	34.73	36.14		
PLACE1002376	3.58	5.91	7.86	6.23	6.82	6.56		
PLACE1002379	6.24	7.66	6.63	10.13	9.68	10.9	**	+
PLACE1002386	0.86	5.32	1.35	1.87	2.05	1.51		
PLACE1002395	3.69	9.97	17.13	16.43	20.62	16.16		
PLACE1002399	2.38	11.09	3.42	5.31	10.38	7.39		
PLACE1002407	1.09	5.22	2.31	2.3	4.01	3.66		
PLACE1002433	1.63	2.17	2.97	2.96	4.35	3.66		
PLACE1002437	0.79	1.4	1.47	1.41	3.28	1.35		
PLACE1002438	0.74	2.38	1.96	1.8	2.43	3.38		
PLACE1002446	4.64	8.42	5.95	10.27	8.88	11.99	*	+
PLACE1002447	1.26	6.06	2.05	3.92	3.14	4.32		
PLACE1002450	1.19	5.92	3.24	4.32	4.21	5.05		
PLACE1002462	0.81	4.02	2.94	3.51	2.99	3.13		
PLACE1002465	0.96	4.69	2.2	2.69	4.31	2.24		
PLACE1002474	1.61	2.26	3.23	3.85	4.41	3.6	*	+
PLACE1002477	11.11	14.51	32.39	44.06	41.42	26.68		
PLACE1002493	3.39	4.1	10.1	14.39	16.66	9.49		
PLACE1002497	0.68	2.81	0.67	1.45	0.93	0.99		
PLACE1002499	2.12	4.73	3	5.98	6.44	5.28	*	+
PLACE1002500	2.61	6.52	7.36	7.58	10.45	7.25		
PLACE1002514	0.3	4.49	1.84	1.74	2.47	1.75		
PLACE1002518	2.86	7.65	6.9	5.62	7.55	4.67		
PLACE1002529	1.14	1.56	2.21	3.19	3.4	1.44		
PLACE1002532	1.31	1.82	3.18	5.75	4.94	5.59	**	+
PLACE1002536	3.59	3.75	3.44	5.84	6.07	3.85		
PLACE1002537	1.63	4.06	2.7	2.69	4.07	3.08		
PLACE1002539	1.86	5.68	2.75	4.53	5.29	4.78		
PLACE1002547	6.09	8.06	7.3	12.32	11.02	11.26	**	+
PLACE1002571	2.84	6.85	5.19	6.84	8.65	6.23		
PLACE1002578	3.57	8.34	8.35	11.11	12.19	8.11		
PLACE1002583	1.33	1.61	2.32	3.18	4.02	2.46		
PLACE1002591	0.82	1.62	2.34	3.25	4.43	1.92		

(1172)

特開2002-191363

2341

2342

PLACE1002598	6.56	10.95	12.39	11.93	9.04	7.74		
PLACE1002604	1.73	3.57	2.69	3.75	5.38	3.51		
PLACE1002612	2.89	8.47	5.95	11.25	10.88	8.06		
PLACE1002625	1.25	4.79	3.18	2.7	3.25	1.82		
PLACE1002638	2.94	8.01	6.66	7.78	6.81	7.29		
PLACE1002655	1.39	6.51	5.57	7.19	7.62	6.46		
PLACE1002665	4.57	3.88	5.4	7.47	12.16	10.75	*	+
PLACE1002685	0.58	1.12	1.3	0.67	2.43	0.98		
PLACE1002692	7.42	8.56	16.7	19.27	22.67	16.29		
PLACE1002714	1.8	3	2.11	2.43	3.14	2.24		
PLACE1002721	2.94	4.37	3.88	5.88	7.1	4.28		
PLACE1002722	0.92	5.42	1.97	1.37	3.28	1.85		
PLACE1002726	1.6	6.24	3.66	4.6	5.7	5.26		
PLACE1002756	1.57	4.5	7.04	5.92	9.63	7.78		
PLACE1002768	1.05	3.72	2.16	2.1	2.34	1.71		
PLACE1002772	0.54	2.15	1.32	2.49	2.86	2.3		
PLACE1002775	4.33	4.71	9.15	7.05	7.08	8.67		
PLACE1002780	185.63	218.72	325.36	272.21	244.38	305.38		
PLACE1002782	0.4	3.76	1.1	1.62	1.69	1.14		
PLACE1002794	1.5	6.71	3.27	2.26	4.59	4.36		
PLACE1002795	1.92	6.45	0.81	2.37	3.63	2.77		
PLACE1002811	0.6	1.57	1.34	1.9	1.46	1.16		
PLACE1002815	6.39	7	10.49	7.24	3.16	9.21		
PLACE1002816	8.5	9.72	9.05	7.22	8.2	7.97	*	-
PLACE1002822	0.58	2.51	2.06	2.2	2.87	1.94		
PLACE1002833	9.98	15.11	16.78	15.42	15.99	16.56		
PLACE1002834	3.2	8.08	6.57	6.23	6.79	8.41		
PLACE1002835	0.62	4.14	1.72	0.85	1.83	1.79		
PLACE1002839	1.13	5.75	2.72	2.89	4.72	2.81		
PLACE1002851	1.52	1.87	1.41	1.98	2.15	2.7		
PLACE1002853	4.18	6.23	9.15	6.26	5.6	7		
PLACE1002881	3.42	5.2	11.04	8.35	11.57	10.26		
PLACE1002901	9.66	12.66	20.09	24.14	25.51	30.67	*	+
PLACE1002904	0.89	7.35	1.41	1.98	1.95	3.09		
PLACE1002905	1.36	5.46	3.26	4.04	4.46	3.4		
PLACE1002908	1.6	5.19	3.18	3.84	5.27	3.81		
PLACE1002911	3.91	6.96	6.9	4.66	7.89	6.75		
PLACE1002941	1.57	2.2	2.48	3.94	2.02	2.31		
PLACE1002950	9.59	9.15	14.74	5.31	8.02	14.51		
PLACE1002955	47.83	40.69	72.7	82.17	62.5	84.64		
PLACE1002958	19.36	26.92	35.27	35.6	35.35	59.02		
PLACE1002962	1.03	4.03	2.2	1.41	2.63	1.67		
PLACE1002967	1.34	4.83	3.19	4.37	3.52	2.81		
PLACE1002968	1.2	5.14	2.7	2.55	3.05	1.81		
PLACE1002976	8.94	12.08	24.23	24.5	36.89	30.05		
PLACE1002991	2.68	3.05	6.66	3.49	4.56	3.6		
PLACE1002993	2.72	3.86	5.52	8.21	6.92	5.56		
PLACE1002996	2.02	3.03	3.43	5.54	3.52	3.01		
PLACE1003010	1.91	3.69	4.27	4.31	3.86	3.32		
PLACE1003025	2.85	7.01	6.1	8.57	11.37	10.11	*	+

(1173)

特開2002-191363

2343

2344

PLACE1003027	5.02	13.08	9.31	8.55	12.45	12.76		
PLACE1003044	1.95	8.24	2.61	3.64	4.16	2.74		
PLACE1003045	1.41	7.75	1.77	1.88	2.64	1.01		
PLACE1003052	2.19	3.16	5.74	4.44	3.6	1.99		
PLACE1003083	1.59	3.04	3.23	3.06	1.61	2.25		
PLACE1003085	3.91	6.19	5.6	9.46	5.89	3.33		
PLACE1003092	3.94	4.87	6.25	7	5.6	6.17		
PLACE1003097	0.37	3.06	1.44	2.12	1.88	1.63		
PLACE1003100	1.65	7.1	4.2	3.88	4.74	4.29		
PLACE1003108	1.26	10.37	2.91	3.32	4.44	2.39		
PLACE1003115	11.39	18.3	58.59	73.64	99.24	69.1	*	+
PLACE1003120	3.1	3.08	9.71	11.34	8.32	10.19		
PLACE1003135	0.72	2.04	1.09	1.56	2.89	1.08		
PLACE1003136	3.95	5.82	6.05	9.03	6.55	7.34		
PLACE1003141	2.04	2.97	2.1	1.97	2.49	1.8		
PLACE1003145	1.21	4.17	2.52	6.24	6.88	7.67	*	+
PLACE1003147	2.87	7.85	5.71	5.02	5.25	6.28		
PLACE1003153	0.54	7.63	2.14	1.66	3.2	1.82		
PLACE1003163	6.09	13.55	8.19	8.39	14.09	12.26		
PLACE1003172	23.21	21.74	44.19	47.78	43.17	39.52		
PLACE1003174	2.31	2.49	3.75	4.3	3.55	1.68		
PLACE1003176	0.47	2	1.89	2.88	1.27	1.46		
PLACE1003181	1.72	4.19	2.72	2.5	2.76	2.36		
PLACE1003184	0.76	3.92	1.53	1.91	1.49	2.06		
PLACE1003190	2.39	9.81	8.67	10.73	7.98	10.34		
PLACE1003200	0.29	4.48	1.84	0.72	1.92	1.16		
PLACE1003205	3.94	7.07	9.68	6.82	10.38	7.2		
PLACE1003209	1.43	2.18	2.62	2.28	1.82	2.89		
PLACE1003214	0.83	1.3	2	2.15	2.44	1.81		
PLACE1003229	2.08	2.78	2.9	2.78	3.35	3.48		
PLACE1003238	0.46	2.34	1.24	1.35	2	0.75		
PLACE1003249	1.87	5.04	4.7	7.33	7.56	5.89		
PLACE1003256	3.47	7.69	7.94	8.82	7.68	6.08		
PLACE1003258	1.03	3.81	3.48	2.42	2.19	0.87		
PLACE1003279	3.09	7.19	9.02	11.15	13.56	11.58	*	+
PLACE1003294	0.95	1.54	1.59	1.57	1.25	2.64		
PLACE1003296	1.49	2.6	2.45	2.59	3.11	2.4		
PLACE1003297	7.52	10.15	31.88	23.01	23.49	19.3		
PLACE1003302	3.92	5.16	6.99	5.8	4.72	5.47		
PLACE1003334	1.51	4.41	1.91	2.4	3.59	3.09		
PLACE1003337	13.69	16.3	29.83	28.53	34.27	25.76		
PLACE1003342	1.05	4.07	1.89	2.15	2.35	1.97		
PLACE1003343	1.07	4.98	1.61	2.02	2.75	2.12		
PLACE1003344	6.25	5.33	12.83	11.18	11.35	11.98		
PLACE1003353	19.67	16.04	40.09	37.78	42.55	40.26		
PLACE1003361	1.82	3.64	3.72	5.85	5.31	4.6	*	+
PLACE1003366	1.45	4.35	3.63	3.22	3.33	2.97		
PLACE1003369	2.75	4.51	3.49	3.29	3.25	4.62		
PLACE1003372	2.08	5.73	2.68	5.45	4.72	3.39		
PLACE1003373	2.85	7.37	6.62	8.8	11.81	9.89		

(1174)

特開 2002-191363

2345

2346

PLACE1003375	1.42	4.91	1.92	2.59	2.95	3.21
PLACE1003378	0.94	0.94	0.98	0.7	1.66	1.08
PLACE1003383	0.87	1.55	2.33	1.59	3.15	1.57
PLACE1003394	10.55	12.49	24.08	11.75	22.99	17.27
PLACE1003401	0.79	3.91	1.34	1.03	2.13	1.04
PLACE1003405	1.5	3.97	2.22	2.54	2.46	2.04
PLACE1003407	2.39	6.06	5.16	3.96	6.3	4.02
PLACE1003420	3.26	7.69	6.19	6.8	10.92	8.7
PLACE1003428	0.63	3.3	2.62	2.07	2.94	1.96
PLACE1003432	6.14	5.81	8.2	6.64	7.05	5.42
PLACE1003438	0.45	2.66	0.93	2.41	2.34	1.99
PLACE1003452	1.87	5.02	5.08	4.53	3.43	3.84
PLACE1003454	2.49	5.59	7.34	7.31	6.95	5.61
PLACE1003455	2.58	4.26	2.35	2.97	3.01	3.17
PLACE1003456	3.22	7.74	8.62	6.9	7.2	7.79
PLACE1003460	6.39	13.35	14.87	13.02	16.76	12.86
PLACE1003478	1.15	1.71	0.86	2.33	2.07	1.24
PLACE1003484	12.06	12.21	45.33	28.12	31.5	34.2
PLACE1003493	1.61	4.72	4.9	3.84	5.96	5.08
PLACE1003503	85.45	87.35	107.79	115.17	111.85	172.81
PLACE1003505	1.99	6.77	4.78	7.44	6.63	8.87
PLACE1003516	0.86	6.78	2.7	2.8	3.95	2.39
PLACE1003519	17.58	26.29	50.41	45.77	36.97	58.75
PLACE1003520	14.18	25.48	35.96	25.73	31.4	32.19
PLACE1003521	2.71	3.64	4.93	5.97	4.71	7.4
PLACE1003525	8.45	11.81	45.05	33.94	43.71	36.88
PLACE1003528	39.18	44.68	136.4	106.04	122.76	127.22
PLACE1003529	1.46	4.26	3.29	2.4	3.94	3.83
PLACE1003537	4.41	9.05	11.05	11.36	13.77	13.3
PLACE1003549	1.1	5.02	4.59	5.61	6.01	5.93
PLACE1003553	1.6	5.89	3.88	4.02	4.17	4.23
PLACE1003566	5.93	9.8	17.51	13.03	19.09	14.58
PLACE1003568	3.01	2.71	5.76	5.69	4.43	4.04
PLACE1003573	0.98	2.43	1.19	2.16	1.87	1.48
PLACE1003575	2.16	3.09	3.44	3.34	3.5	4.15
PLACE1003583	0.97	3.45	3.34	2.29	4.16	2.23
PLACE1003584	1.23	4.46	4.01	3.56	3.8	2.65
PLACE1003592	4.4	8.48	9.48	7.11	10.48	9.53
PLACE1003593	0.84	5.55	2.4	2.23	3.02	1.55
PLACE1003594	4.24	6.76	6.01	5.78	6.68	7.36
PLACE1003596	13.77	11.31	22.53	15.68	17.69	8.25
PLACE1003598	2.83	3.63	5.02	4.89	3.7	2.95
PLACE1003602	1.8	4.24	6.36	4.37	3.54	2.68
PLACE1003605	17.43	21.72	45.86	94.65	95.6	91.55
PLACE1003611	2.34	5.18	6.07	5.65	7.14	6
PLACE1003618	0.67	7.39	2.09	1.58	2.4	1.32
PLACE1003625	1.78	10.41	2.75	3.33	5.48	2.8
PLACE1003626	8.77	15.99	17.14	10.87	14.46	13.19
PLACE1003630	1.8	2.57	5.8	7.05	4.67	5.86
PLACE1003635	2.15	1.83	3.19	2.96	2.82	2.44

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(1175)

特開2002-191363

2347

2348

PLACE1003638	1.3	2.58	3	5.21	4.19	3.13		
PLACE1003644	4.01	5.7	7.25	7.81	9.13	8.76	*	+
PLACE1003654	2.56	6.14	4.04	3.54	6.96	6.4		
PLACE1003656	2.69	7.79	6.12	6.54	5.63	4.77		
PLACE1003660	0.26	9.54	3.5	3.08	4.92	4.11		
PLACE1003669	2.43	9.05	3.67	2.59	4.26	3.08		
PLACE1003670	5.37	5.7	9.44	11.01	8.26	8.76		
PLACE1003671	1.66	1.22	3.57	3.11	2.57	1.57		
PLACE1003697	7.27	7.99	9.8	8.23	6.06	6.42		
PLACE1003704	3.12	3.97	5.17	5.96	7.25	5.97	*	+
PLACE1003709	0.89	2.63	0.8	1.19	1.24	2.44		
PLACE1003711	0.74	5.48	1.35	1.87	1.8	1.39		
PLACE1003723	1.07	6.99	4.7	4.2	5.31	4.16		
PLACE1003724	3.31	10.74	9.1	9.11	11.79	10.49		
PLACE1003737	2.14	2.21	4.72	3.35	3.1	3.29		
PLACE1003738	1.06	1.94	3.13	3.96	3.92	3.41		
PLACE1003742	2.25	3.58	5.71	6.81	6.18	2.85		
PLACE1003744	6.13	8.86	14.6	16.21	17.96	19.13	*	+
PLACE1003758	0.85	4.55	0.96	2	1.46	1.16		
PLACE1003760	13.44	18.68	27.23	31.82	20.52	22.79		
PLACE1003762	1.45	4.97	3.7	3.77	3.78	3.49		
PLACE1003765	1.18	5.23	3.45	2.01	3.1	2.11		
PLACE1003768	0.36	1.14	1.36	1.45	2.75	1.07		
PLACE1003771	1.28	1.94	2.07	1.84	3.13	1.43		
PLACE1003772	34.15	38.19	97.86	62.42	64.06	52.43		
PLACE1003783	1.48	3.02	2.22	18.65	19.53	16.61	**	+
PLACE1003784	0.69	3.92	0.87	2.09	2.19	2.68		
PLACE1003788	0.4	4.92	1.06	1.85	1.71	0.32		
PLACE1003795	1.01	4.1	3.57	4.73	4.38	3.54		
PLACE1003827	13.83	20.46	20.72	22.48	30.84	25.92		
PLACE1003833	0.98	1.49	3.9	3.65	4.33	3.31		
PLACE1003839	22.55	19.18	52.95	50.39	56.11	43.86		
PLACE1003845	6.09	6.88	11.72	24.98	19.99	10.6		
PLACE1003850	3.16	4.84	7.19	5.45	5.95	6.39		
PLACE1003852	0.25	3.36	1.09	0.99	1.58	1.05		
PLACE1003858	1.34	4.42	1.99	2.04	1.85	2.92		
PLACE1003861	0.95	4.51	1.63	2.98	2.78	1.73		
PLACE1003864	0.94	5.5	2.74	2.88	3.86	2.69		
PLACE1003870	3.84	3.4	13.2	10.71	16.12	12.03		
PLACE1003885	1.33	1.42	1.59	3.07	3.76	2.38	*	+
PLACE1003886	4.56	6.01	5.75	9.27	6.87	4.3		
PLACE1003888	0.75	3.79	1.96	2.87	3.42	2.68		
PLACE1003892	4.93	6.91	20.79	17.21	22.33	14.87		
PLACE1003900	2.27	5.92	7.17	6.34	9.75	6.6		
PLACE1003902	1.91	7.39	5.25	5.43	8.07	6.29		
PLACE1003903	0.42	5.07	2.94	2.55	3.5	2.59		
PLACE1003915	8.15	7.04	10.78	8.31	9.79	9.84		
PLACE1003918	1.88	2.45	4.75	3.47	6.26	3.75		
PLACE1003923	2.06	3.73	5.63	2.7	5.54	3.32		
PLACE1003932	3.99	5.16	5.47	4.06	7.58	5.09		

(1176)

特開2002-191363

2349

2350

PLACE1003936	1.02	3.82	2.81	3.63	2.42	2.78		
PLACE1003966	3.11	7.43	7.76	4.89	7.32	4.21		
PLACE1003968	1.68	5.68	5.94	3.33	4.26	4.57		
PLACE1004018	25.49	33.73	48.16	32.56	40.53	28.62		
PLACE1004020	8.91	10.18	13.26	11.42	18.03	24.15		
PLACE1004028	0.41	2.55	1.3	1.38	1.23	1.91		
PLACE1004034	3.56	4.53	5.22	8.42	9.03	13.04	*	+
PLACE1004042	17.25	20.19	68.35	79.51	93.32	79.44	*	+
PLACE1004078	1.14	4.1	3.3	4.95	6.51	4.45		
PLACE1004103	5.54	10.93	13.98	13.77	15.09	14.14		
PLACE1004104	5.94	12.29	27.78	24.18	34.98	29.71		
PLACE1004113	1.37	3.7	3.28	2.75	3.27	1.29		
PLACE1004114	1.12	2.55	2.23	2.84	2.49	2.93		
PLACE1004118	1.58	3.52	2.09	2.74	3.64	2.5		
PLACE1004128	4.4	6.84	5.66	6.61	8.13	9.01		
PLACE1004130	2.25	4.83	8.35	6.86	9.26	6.73		
PLACE1004149	3.59	7.3	9.86	10.23	13.29	10.12		
PLACE1004156	3.61	7.91	10.12	11.66	17.62	13.31		
PLACE1004160	5.45	9.54	14.36	13.1	19.65	14.53		
PLACE1004161	2.2	4.86	3.54	4.85	5.73	7.48		
PLACE1004166	5.61	5.81	9.91	8.3	8.66	11.2		
PLACE1004168	3.35	4.97	3.73	4.73	6.65	6.79		
PLACE1004170	0.78	3.28	1.93	2.98	3.42	2.76		
PLACE1004178	0.83	5.23	2.37	2.4	2.59	2.36		
PLACE1004183	0.89	7.99	4.41	3.53	4.32	4.84		
PLACE1004197	0.64	5.14	1.55	1.73	3.54	1.65		
PLACE1004199	1.66	4.52	4.09	3.78	5.88	4.35		
PLACE1004203	1.8	3.57	4.17	2.43	3.62	2.83		
PLACE1004242	3.8	5.64	11.04	8.55	8.14	8.64		
PLACE1004249	31.4	56.31	117.88	127.93	152.54	151.22	*	+
PLACE1004255	0.79	2.65	1.26	2.59	2.15	1.93		
PLACE1004256	9.06	11.68	13.63	14.66	14.18	23.37		
PLACE1004257	2.63	7.95	6.48	7.89	8.8	8.64		
PLACE1004258	1.87	5.21	3.13	4.59	3.15	3.11		
PLACE1004270	0.72	3.8	2.5	2.7	4.01	1.65		
PLACE1004272	1.34	3.68	3.73	3.86	5.38	6.15		
PLACE1004273	92.91	89.59	212.62	212.05	129.56	99.82		
PLACE1004274	2.09	3.61	6.51	6.42	7.14	6.74		
PLACE1004277	2.3	4.4	5.76	6.45	7.7	6.04		
PLACE1004279	0.54	3.39	2.23	3.16	2.64	2.02		
PLACE1004282	2.43	8.25	6.62	4.22	5.56	4.49		
PLACE1004284	4.59	11.31	7.84	7.38	8.16	7.15		
PLACE1004289	1.28	7.85	2.46	3.06	3.63	2.1		
PLACE1004299	0.33	6.41	1.38	1.54	2.67	1.83		
PLACE1004302	1.01	2.98	3.27	2.41	2.45	1.12		
PLACE1004305	1.11	2.09	1.9	1.9	1.82	1.78		
PLACE1004316	2.3	4.48	5.4	6.06	3.85	4.52		
PLACE1004322	2.49	3.41	5.25	6.35	7.14	5.75	*	+
PLACE1004325	2.43	6.38	3.84	3.85	3.66	4		
PLACE1004332	1.21	7.18	2.8	3.72	3.46	3.01		

(1177)

特開2002-191363

2351

2352

PLACE1004336	2.87	9.6	5.36	6.88	9.21	6.71		
PLACE1004346	0.47	7.22	2.58	1.87	2.54	1.69		
PLACE1004358	1.3	2.41	2.41	2.46	2.09	2.03		
PLACE1004376	11.07	10.15	23.35	19.16	17.04	16.46		
PLACE1004384	0.65	3.46	1.46	1.92	2.48	2.1		
PLACE1004385	1.4	2.89	1.69	3.52	1.88	2.67		
PLACE1004388	1.79	5.73	4.27	3.44	4.87	3.04		
PLACE1004405	2.16	8.39	4.42	13.36	14.48	16.74	**	+
PLACE1004407	5.05	13.12	13.37	11.2	16.24	11.85		
PLACE1004424	0.37	5.78	0.85	1.7	1.81	1.58		
PLACE1004425	1.14	1.94	3.57	3.28	3.27	3.44		
PLACE1004427	1.96	3.31	4.56	4.67	4.22	3.24		
PLACE1004428	0.88	2.05	2.17	2.66	2.08	2.62		
PLACE1004433	5.7	8.3	10.82	12.94	15.67	12.05	*	+
PLACE1004435	0.72	4.17	1.43	1.95	1.9	2.15		
PLACE1004437	4.05	7.68	14.2	11.07	13.01	12.37		
PLACE1004441	7.82	11.68	34.06	30.75	43.19	26.41		
PLACE1004446	1.5	4.36	0.9	1.03	1.35	1.39		
PLACE1004450	0.33	1.46	1.34	2.57	1.71	0.7		
PLACE1004451	0.51	1.45	2.14	1.89	2.69	0.88		
PLACE1004456	8.22	9.7	10.97	16.68	10.4	4.18		
PLACE1004458	3.39	4.81	3.66	7.77	7.05	8.24	**	+
PLACE1004460	0.84	4.58	2.1	2.91	2.69	1.75		
PLACE1004467	5.31	6.81	10.65	7.67	10.14	10.48		
PLACE1004471	2.65	5.93	6.64	6.79	7.34	6.14		
PLACE1004473	1.16	4.66	3.5	3.18	3.23	3.21		
PLACE1004475	14.03	16.41	32.49	31.09	32.51	18.17		
PLACE1004482	8.37	6.7	10.79	10.04	9.76	9.15		
PLACE1004491	0.39	2.51	1.49	1.19	2.68	1.3		
PLACE1004492	61.52	74.8	127.94	129.92	127.64	123.82		
PLACE1004506	10.71	14.35	14.4	8.45	11.13	10.03		
PLACE1004507	2.9	7.37	5.09	7.15	6.87	6.18		
PLACE1004510	2.51	6.23	6.33	6.59	7.8	8.16		
PLACE1004516	0.98	7.36	2.12	2.79	3.78	2.22		
PLACE1004518	1.64	1.78	3.03	2.41	3.88	2.83		
PLACE1004519	0.17	0.82	0.62	1.43	2.79	1.51	*	+
PLACE1004520	6.08	8.09	10.06	7.44	9.11	2.52		
PLACE1004530	33.19	43.86	68.13	41.86	27.72	38.09		
PLACE1004545	1.13	3.83	2.12	3.03	3.31	3.65		
PLACE1004547	5	7.61	7.82	8.66	11.2	10.28		
PLACE1004548	1.69	6.73	4.43	6.93	8.48	6.5		
PLACE1004550	2.27	6.24	6.67	5.92	6.78	6.15		
PLACE1004551	0.8	2.16	1.62	2.14	2.21	1.95		
PLACE1004559	2.9	2.89	5.11	4.45	6.75	4.82		
PLACE1004562	8.67	11.27	16.07	13.01	14.38	13.34		
PLACE1004564	1.84	5.19	4.36	4.64	5.98	4.49		
PLACE1004604	1.69	4.21	9.88	2.49	4.34	1.97		
PLACE1004611	2.73	5.87	4.89	3.86	4.17	3.99		
PLACE1004629	9.42	15.75	19.92	23.93	30.49	29.01	*	+
PLACE1004630	16.66	20.82	35.1	16.76	23.04	19.17		

(1178)

特開2002-191363

2353

2354

PLACE1004637	5.03	8.82	10.34	6.61	9.17	8.29
PLACE1004645	36.5	39.28	92.04	85.16	87.94	74.59
PLACE1004646	1.07	2.91	2.87	3.68	2.19	2.28
PLACE1004648	0.8	3.42	2.52	2.53	3.15	1.22
PLACE1004655	45.95	58.09	130.94	112.14	126.25	99.01
PLACE1004658	2.4	7.34	6.31	6.64	8.37	6
PLACE1004664	1.26	4.83	1.3	3.02	2.65	1.89
PLACE1004672	2.32	6.79	8.02	6.3	7.51	7.14
PLACE1004674	9.4	11.97	14.7	8.3	9.75	15.25
PLACE1004681	1.97	3.84	5.62	5.05	5.69	4.39
PLACE1004686	2.74	4.33	7.46	9.22	10.77	8.59
PLACE1004690	10.64	13.43	19.62	21.75	19.21	31.72
PLACE1004691	1.14	6.71	3.71	2.92	4.13	2.75
PLACE1004693	1.34	7.54	4.89	3.91	4.59	5.97
PLACE1004701	13.01	18.45	24.24	25.21	24.46	25.1
PLACE1004705	1.29	3.33	2.27	1.8	1.96	1.47
PLACE1004708	37.69	46.37	80.19	41.34	39.66	50.98
PLACE1004716	6.37	8.81	11.08	4.22	12.55	14.26
PLACE1004722	1.31	3.05	2.6	2.26	3.28	2.51
PLACE1004736	5.25	7.71	7.6	9.16	8.89	11.63
PLACE1004737	5.42	12.71	16.14	8.15	11.23	13.78
PLACE1004740	4.88	9.06	8.22	7.37	7.93	8.2
PLACE1004743	1.31	4.04	3.1	1.97	4	3.55
PLACE1004751	0.98	2.89	2.88	2.75	3.74	3.06
PLACE1004757	3.45	4.34	10.53	8.4	9.6	7.22
PLACE1004761	6.41	7.32	12.59	9.99	10.44	9.72
PLACE1004773	1.05	2.34	1.7	1.94	2.31	2.72
PLACE1004775	0.35	3.26	1.37	1.29	2.14	1.07
PLACE1004777	2.1	7.57	2.97	3.68	4.25	4.4
PLACE1004793	0.83	4.58	1.37	1.9	2.06	1.25
PLACE1004796	6.65	8.7	13.08	7.79	8.57	7.88
PLACE1004804	0.99	4.46	3.25	2.44	2.38	3.22
PLACE1004813	4.55	7.11	9.84	6.45	5.19	5.45
PLACE1004814	7.16	11.76	17.62	15.83	11.39	10.1
PLACE1004815	0.7	2.81	2.43	3.12	2.61	3.44
PLACE1004816	1.16	2.63	2.04	2.36	2.26	1.84
PLACE1004824	3.25	7.37	5.27	8.1	9.13	8.85
PLACE1004827	1.4	10.89	3.17	2.57	3.05	1.3
PLACE1004836	1.72	12.95	4.26	6.25	7.99	4.49
PLACE1004838	1.35	8.81	2.2	2.49	2.34	1.68
PLACE1004840	1.59	2.06	2.21	1.85	2.08	1.37
PLACE1004842	0.86	1.98	1.89	1.98	1.78	2.33
PLACE1004850	0.81	2.35	1.63	1.83	2.76	2.36
PLACE1004868	0.81	2.97	2.04	1.62	2.23	2
PLACE1004885	1.5	7.09	3.51	3.4	6.03	4.77
PLACE1004886	1.87	8.53	2.76	3.33	5.12	4.93
PLACE1004887	18.14	34.01	58.51	36.9	38.66	30.33
PLACE1004896	8.39	14.15	15.4	9.39	11.14	14.03
PLACE1004900	1.75	2.7	6.69	5.66	5.74	4.44
PLACE1004902	5.42	6.25	9.27	6.39	4.2	3.6

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(1179)

特開2002-191363

2355

2356

PLACE1004904	1.7	4.66	2.52	6.49	4.78	3.03		
PLACE1004911	0.69	2.5	1.12	5.95	4.82	6.09	**	+
PLACE1004913	3.63	5.72	7.38	4.49	5.45	4.79		
PLACE1004918	1.3	6.69	2.05	2.39	3.24	2.64		
PLACE1004930	2.74	8.84	5.93	10.63	16.57	14.71	*	+
PLACE1004934	1.14	4.3	3.34	2.97	2.03	2.5		
PLACE1004937	2.1	4.03	4.91	2.74	3.59	2.36		
PLACE1004949	4.32	4.98	7.67	8.53	8.48	6.04		
PLACE1004969	0.74	1.74	1.99	1.39	2.34	1.4		
PLACE1004970	0.45	2.18	1.2	1.43	1.31	1.01		
PLACE1004972	1.63	6.56	2.69	5.66	3.17	4.76		
PLACE1004974	1.27	5.21	4.2	5.6	5.7	5.62		
PLACE1004975	0.59	2.84	1.11	1.94	1.98	1.24		
PLACE1004979	1.58	4.06	4.26	4.91	5.69	4.75		
PLACE1004982	5.66	6.45	9.74	10.03	10.67	5.65		
PLACE1004985	1.4	1.47	1.46	2.5	2.17	1.2		
PLACE1005003	2.85	4.22	6	6.05	6.37	5.78		
PLACE1005004	0.47	3.36	0.92	1.5	0.73	0.85		
PLACE1005005	3.35	6.9	5.32	6.67	9.65	7.17		
PLACE1005011	6.03	11.12	35.8	37.66	56.97	36.62		
PLACE1005026	0.79	4.18	2.29	2.44	2.49	2.16		
PLACE1005027	2.46	6.72	6.69	5.36	6.92	5.68		
PLACE1005031	1.21	1.47	3.69	5.72	7.1	4.6	*	+
PLACE1005036	2.27	3.6	6.83	9.08	8.91	7.53	*	+
PLACE1005041	2.5	2.84	2.84	5.05	4.41	3.25	*	+
PLACE1005046	2.23	4.11	4.56	6.04	3.92	4.33		
PLACE1005047	0.23	3.19	2.6	1.43	1.23	2.1		
PLACE1005052	4.24	8.53	6.36	8.08	8.15	8.1		
PLACE1005055	2.54	7.45	4.66	7.2	6.45	5.62		
PLACE1005066	4.33	8.26	7.58	12.9	14.14	16.49	**	+
PLACE1005077	1.17	0.68	1.2	2.1	2.43	1.54	*	+
PLACE1005085	1.41	1.97	3.06	3.34	4.14	3.45		
PLACE1005086	1.93	3.77	5.17	5.62	7.78	4.79		
PLACE1005088	24.66	32.47	46.03	43.45	31.47	27.46		
PLACE1005089	1.57	4.78	3.15	2.52	3.67	3.14		
PLACE1005101	3.37	8.11	5.46	6.11	8.96	6.39		
PLACE1005102	2.56	7.14	5.01	4.11	5.51	3.8		
PLACE1005108	2	6.08	5.87	6.58	6.19	5.09		
PLACE1005110	1.34	1.89	3.08	1.75	2.75	2.15		
PLACE1005111	1.31	1.34	1.23	1.45	2.17	1.54		
PLACE1005123	26.23	26.21	47.58	34.26	49.34	34.98		
PLACE1005124	3.2	4.66	4.18	4.2	6.91	5.44		
PLACE1005128	9.54	8.89	18.22	16.37	16.36	16.13		
PLACE1005130	2.65	6.57	5.54	2.84	4.98	3.58		
PLACE1005141	6.3	9.92	11.25	16.15	20.75	18.95	**	+
PLACE1005146	1.3	2.71	3.03	2.17	2.53	2.29		
PLACE1005152	1.85	3.9	4.56	4.13	4.23	4.79		
PLACE1005157	2.66	5.19	5.3	4.38	4.09	7.01		
PLACE1005162	2.79	3.72	9.31	6.57	7.45	7.1		
PLACE1005170	17.34	18.92	29.76	21.38	18.18	23.73		

(1180)

特開2002-191363

2357

2358

PLACE1005176	0.57	5.6	1.7	2.33	2.47	1.94		
PLACE1005181	0.53	5.14	0.96	0.89	1.36	0.37		
PLACE1005184	4.06	9.09	10.4	8.97	12.82	11.26		
PLACE1005186	3.5	3.41	8.56	8.05	5.79	5.73		
PLACE1005187	2.85	4	4.13	6.1	4.99	4.25		
PLACE1005189	6.12	7.71	5.34	10.84	10.65	12.22	**	+
PLACE1005193	1.48	3.78	1.71	3.84	2.91	2.61		
PLACE1005200	1.35	4.68	2.61	2.47	3.75	3.1		
PLACE1005206	2.43	6.48	4.26	3.35	3.95	2.95		
PLACE1005216	1.53	5.46	4.44	5.6	6.51	4.12		
PLACE1005223	1.43	6.21	5	4.38	5.66	3.27		
PLACE1005225	1.36	3.01	3.49	3.33	3.32	4.65		
PLACE1005232	1.86	3.31	4.87	5.63	6.19	3.88		
PLACE1005239	1.06	4.3	2.32	2.84	2.86	2.41		
PLACE1005243	4.35	7.32	5.41	8.48	7.49	10.75		
PLACE1005250	4.24	10.31	7.98	4.38	5.9	8.88		
PLACE1005261	3.21	7.43	4.74	4.78	5.82	3.51		
PLACE1005266	1.05	4.47	2.82	2.28	4.43	2.76		
PLACE1005271	4.66	5.31	8.79	5.87	11.16	7.95		
PLACE1005277	2.06	3.48	2.35	2.62	1.98	2.64		
PLACE1005287	3.63	4.31	5.87	2.98	5.06	6.91		
PLACE1005299	24.16	22.75	48.29	35.17	24.24	41.06		
PLACE1005305	6.81	8.46	11.13	10.67	11.85	16.25		
PLACE1005307	1.59	5.44	4.14	3.15	5.42	4.84		
PLACE1005308	2.41	4.96	3.95	5.32	5.99	5.79		
PLACE1005313	1.08	3.83	1.6	1.8	2.05	1.8		
PLACE1005320	1.36	3.65	3.34	3.39	4.05	2.26		
PLACE1005327	10.78	8.74	16.8	10.36	7.95	4.43		
PLACE1005331	2.28	4.92	5.28	4.66	4.97	3.33		
PLACE1005335	1.53	3.8	2.24	2.03	3.22	2.42		
PLACE1005336	9.12	12.58	16.58	16.39	16.99	20.15		
PLACE1005351	2.62	8.18	10.17	9.28	8.66	9.52		
PLACE1005366	2.04	6.93	3	2.99	3.71	4.23		
PLACE1005373	1.77	6.34	4.44	3.91	5.36	3.37		
PLACE1005374	3.29	9.47	11.4	7.35	10.22	12.41		
PLACE1005383	8.16	7.54	12.81	7.21	5.93	4.03		
PLACE1005388	0.33	2.04	1.56	1.92	3.67	2.2		
PLACE1005409	2.97	5.02	4.99	3.9	4.23	2.97		
PLACE1005410	12.41	16.44	18.89	24.38	20.98	27.1	*	+
PLACE1005426	5.16	7.48	9.06	5.51	7.67	5.45		
PLACE1005431	12.6	15.65	22.53	19.64	26.25	23.75		
PLACE1005453	1.4	10.38	3.93	4.85	4.45	3.28		
PLACE1005467	3.09	11.87	7	5.57	11.63	7.28		
PLACE1005471	1.6	1.94	1.66	2.29	1.52	1.28		
PLACE1005476	0.42	1.73	1.24	1.6	1.57	1.46		
PLACE1005477	1.58	2.26	2.51	3	2.93	2.74		
PLACE1005480	0.77	2.01	1.86	1.93	1.4	0.53		
PLACE1005481	0.44	4.81	2.3	2.77	3.62	2.44		
PLACE1005494	0.27	6.66	1.68	1.21	1.73	1.06		
PLACE1005495	3.86	12.83	8.31	6.85	9.25	7.62		

(1181)

特開2002-191363

2359

2360

PLACE1005497	2.27	7.72	3.95	4.24	5.68	5.91		
PLACE1005499	5.71	5.86	11.07	10.82	7.9	6.49		
PLACE1005502	1.59	2.87	3.43	4.07	3.45	1.49		
PLACE1005513	1.77	4.14	3.35	1.86	2.85	1.98		
PLACE1005515	2.89	4.76	4.22	4.58	5.29	3.78		
PLACE1005519	1.04	4.53	3.29	2.85	2.85	2.83		
PLACE1005526	0.58	5.55	1.38	1.3	1.59	0.71		
PLACE1005528	2.08	7.71	5.57	5.94	7.12	5.33		
PLACE1005530	2.16	7.09	4.32	5.17	8.23	4.67		
PLACE1005536	1.74	1	2.74	3.12	2.43	2.88		
PLACE1005539	10.1	11.64	23.77	8.65	8.66	5.22		
PLACE1005543	1.7	3.57	5.62	3.54	4.32	2.57		
PLACE1005544	0.86	3.26	3.15	2.49	2.68	2.27		
PLACE1005550	4.32	7.61	7.85	10.16	7.25	6.86		
PLACE1005554	1.15	5.47	2.67	2.17	2.17	1.17		
PLACE1005557	1.76	7.21	4.95	8.22	7.64	7.7		
PLACE1005563	0.51	4	1.89	1.45	2.07	1.06		
PLACE1005569	0.6	0.5	1.56	1.59	1.81	1.09		
PLACE1005574	1.07	1.88	2.49	2.48	4.43	2.22		
PLACE1005584	1.3	2.68	3.91	3.91	5.58	3.03		
PLACE1005590	4.28	5.14	8.4	9.87	10.73	8.02		
PLACE1005595	3.08	4.03	2.89	3.65	3.81	3.89		
PLACE1005601	2	5.66	4.22	3.77	4	4.02		
PLACE1005603	1.08	4.9	1.04	2.49	0.95	1.94		
PLACE1005604	1.2	6.71	2.42	3.6	4.2	3.46		
PLACE1005611	2.22	2.3	3.98	5.15	5.65	2.89		
PLACE1005622	0.65	1.71	1.98	2.94	3.88	1.26		
PLACE1005623	1.42	3.08	3.27	3.71	3.65	1.61		
PLACE1005630	3.31	5.81	7.75	87.83	72.15	89.12	**	+
PLACE1005639	0.75	4.36	1.28	1.66	2.02	1.18		
PLACE1005646	2.13	5.41	4.31	5.4	5.08	2.55		
PLACE1005647	2.77	9.69	6.72	7.34	9.11	6.25		
PLACE1005648	3	8.11	9.21	8.34	10.59	8.22		
PLACE1005653	1.99	1.43	2.74	2.74	2.13	2.67		
PLACE1005656	0.92	2.48	2.24	1.68	2.78	1.58		
PLACE1005659	0.87	2.64	1.01	1.62	1.84	1.32		
PLACE1005660	3.91	8.03	5.77	8.87	8.88	8.34		
PLACE1005664	2.69	6.57	6.14	3.39	4.27	3.19		
PLACE1005666	0.89	5.91	3.55	4.63	4.93	3.97		
PLACE1005669	4.46	10.41	11.39	11.64	13.9	14.6		
PLACE1005682	1.94	5.27	4.49	6.2	5.2	5.47		
PLACE1005698	0.6	2.7	2.92	2.01	2.67	2.38		
PLACE1005708	25.32	34.08	53.46	53.89	59.98	53.76		
PLACE1005725	3.25	3.75	6.41	5.64	5.82	7.29		
PLACE1005727	2.97	4.54	4.15	3.9	3.49	4		
PLACE1005730	0.77	4.29	3.26	1.1	1.54	1.28		
PLACE1005736	5.37	7.55	5.73	9.25	12.55	10.19	*	+
PLACE1005739	0.81	4.96	1.38	2.46	3.17	1.74		
PLACE1005745	8.03	7.11	11.52	11.98	6.97	11.44		
PLACE1005752	1.31	3.15	2.96	2.55	2.24	1.25		

(1182)

特開2002-191363

2361

2362

PLACE1005755	0.8	2.79	3.02	1.72	3.28	2.27		
PLACE1005756	10.79	12.06	17.2	18.22	19.3	21.47	*	+
PLACE1005760	10.22	15.24	68.06	49.69	68.81	53.09		
PLACE1005763	1.47	7.04	3.58	3.79	4.63	3.02		
PLACE1005768	1.25	5.63	3.69	4.58	5.13	4.19		
PLACE1005771	5.71	13.63	13.7	11.28	17.49	17.27		
PLACE1005783	1.82	2.44	3.64	3.05	3.71	3.47		
PLACE1005799	4.79	5.25	8.37	6.12	8.78	8.62		
PLACE1005802	1.07	3.78	3.64	2.7	3.64	1.96		
PLACE1005803	3.06	6.15	4.78	5.6	4.94	7.36		
PLACE1005804	0.92	8.41	1.33	2	1.91	2.44		
PLACE1005813	17.23	18.71	78.06	70.01	94.17	74.89		
PLACE1005815	1.43	5.6	4.38	3.8	5.01	4.1		
PLACE1005828	2.11	3.62	4.42	5.34	6.24	3.56		
PLACE1005833	119.17	92.82	182.22	122	114.37	107.96		
PLACE1005834	2.04	4.33	3.95	3.55	3.56	2.56		
PLACE1005835	22.7	19.1	51.52	72.32	60.34	68.56	*	+
PLACE1005836	2.39	4.21	4.97	2.61	3.83	2.55		
PLACE1005845	0.97	5.42	2.66	2.67	3.05	3.65		
PLACE1005850	1.82	3.91	3.04	2.84	2.85	2.15		
PLACE1005851	1.03	3.44	1.46	1.2	2.01	1.11		
PLACE1005856	0.92	4.01	2.42	2.24	3.37	3.37		
PLACE1005875	1.78	3.89	4.77	3.3	3.48	3.17		
PLACE1005876	1.33	3.99	4.76	6.87	6.34	6.9	*	+
PLACE1005878	1.3	2.67	2.08	3.54	4.46	2.79		
PLACE1005880	2.36	4.09	4.31	4.16	3.07	3.45		
PLACE1005884	1.6	4.87	1.89	2.48	2.21	2.73		
PLACE1005890	1.9	9.57	3.7	2.26	3.09	1.88		
PLACE1005898	3.29	10.87	5.34	7.36	8.12	6.36		
PLACE1005913	1.46	9.31	8.05	4.99	6.47	4.33		
PLACE1005921	0.99	1.92	2.03	1.49	2.22	0.79		
PLACE1005923	0.74	1.61	1.17	1.47	2.42	1.43		
PLACE1005925	0.83	2.67	3.18	2.19	2.14	1.68		
PLACE1005927	1.26	2.49	1.93	1.95	2.56	2.3		
PLACE1005932	2.04	5.66	2.44	2.53	2.32	2.52		
PLACE1005934	0.88	7.91	3.16	3.9	5.61	4.19		
PLACE1005936	1.31	8.96	3.02	2.02	2.84	2.3		
PLACE1005939	54.61	68.58	111.22	157.61	194.58	212.18	**	+
PLACE1005951	2.36	3.39	4.98	5.56	4.48	2.35		
PLACE1005953	1.5	1.64	2.64	2.59	2.43	3.03		
PLACE1005955	1.64	2.01	3.8	4.07	3.43	2.55		
PLACE1005966	0.76	3.42	1.69	1.75	2	2.19		
PLACE1005968	1.52	4.96	3.2	4.71	5.15	6.12		
PLACE1005975	2.58	7.11	5.42	6.18	7.01	6.49		
PLACE1005990	0.7	7.7	1.54	2.1	1.87	0.88		
PLACE1005997	88.15	118.52	196.48	189.6	226.97	172.1		
PLACE1006002	3.38	3.97	8.87	8.4	7.71	9.18		
PLACE1006003	1.55	3.02	4.83	5.09	4.44	5.33		
PLACE1006011	1.85	3.63	3.46	4.48	2.68	1.91		
PLACE1006017	0.84	2.74	2.81	3.4	3.4	3.58		

(1183)

特開 2002-191363

2363

2364

PLACE1006037	2.99	7.05	2.48	6.14	3.64	4.29		
PLACE1006040	2.2	7.87	3.97	6.64	6.9	7.77		
PLACE1006063	0.94	4.64	2.59	2.11	3.15	2.25		
PLACE1006071	3.06	6.52	4.97	5.36	4.03	4.47		
PLACE1006073	2.74	3.53	6.43	7.19	6.81	6.93		
PLACE1006074	1.4	2.22	3.34	2.62	3.23	1.69		
PLACE1006076	1.36	2.51	2.98	3.15	2.47	2.75		
PLACE1006079	1.38	4.32	1.78	2.1	1.1	1.11		
PLACE1006093	0.49	3.76	1	3.56	3.85	1.83		
PLACE1006116	2.99	6.44	4.04	5.28	5.01	4.91		
PLACE1006119	3.15	6.81	7.07	9.22	10.4	8.03		
PLACE1006129	2.12	5.6	3.98	6.59	7.62	5.65		
PLACE1006139	3.44	2.98	6.03	7.77	8.85	5.58		
PLACE1006143	0.5	1.48	1.87	3.18	4.13	3.17	*	+
PLACE1006157	1.55	2.54	4.82	2.96	3.9	2.44		
PLACE1006159	0.69	3.61	0.94	2.68	1.98	1.04		
PLACE1006164	0.35	3.18	1.37	1.73	1.85	1.21		
PLACE1006167	2.18	6.5	3.37	3.95	4.52	3.13		
PLACE1006170	2.79	6.09	6.09	4.34	5.31	3.68		
PLACE1006181	2.75	7.34	2.84	5.8	5.51	5.22		
PLACE1006187	0.76	1.3	2.15	2.01	2.41	1.48		
PLACE1006195	0.11	1.24	1.73	1.93	1.93	0.87		
PLACE1006196	1.8	4.01	4.15	4.32	5.77	2.19		
PLACE1006197	2.12	5.6	5.24	4	3.47	3.39		
PLACE1006198	0.27	3.68	1.21	0.84	1.63	0.5		
PLACE1006205	0.89	5.59	0.99	2.43	2.18	1.28		
PLACE1006208	7.28	13.32	13.46	14.09	14.99	12.5		
PLACE1006211	2.6	8.05	7.92	6.07	9.08	9.35		
PLACE1006219	6.77	5.77	8.94	14.88	22.25	15.35	*	+
PLACE1006223	1.55	1.46	3.19	1.39	3	1.64		
PLACE1006225	0.56	2.27	1.3	1.04	2.3	0.99		
PLACE1006236	1.53	3.2	2.92	3.06	5.01	2.29		
PLACE1006239	0.67	3.62	1.97	2.61	3.66	3.41		
PLACE1006245	3.86	7.13	5.45	4.43	7.44	3.28		
PLACE1006246	1.66	6.56	6.19	5.59	7.66	6.33		
PLACE1006248	1.58	4.47	5.6	2.77	3.1	2.82		
PLACE1006262	0.93	2.24	1.49	2.08	1.61	1.4		
PLACE1006269	2.28	4.71	3.42	2.06	2.47	2.33		
PLACE1006275	1.6	3.57	3.37	4.12	3.68	3.53		
PLACE1006277	1.01	2.42	1.4	1.79	3.01	0.88		
PLACE1006288	9.32	13.59	22.49	26.85	18.4	25.21		
PLACE1006290	1.79	6.81	5.99	8.87	7.56	9.13		
PLACE1006298	1.93	5.52	2.47	3.87	5.08	4.55		
PLACE1006311	0.65	3.38	1.75	225.97	161.43	251.12	**	+
PLACE1006318	3.52	4.03	4.17	4.04	3.17	4.01		
PLACE1006325	5.43	6.73	6.31	8.09	8.38	8.08	**	+
PLACE1006331	1.87	3.36	3.21	4.44	3.59	2.56		
PLACE1006335	1.76	3.64	2.55	4.45	2.98	2.92		
PLACE1006357	0.27	4.51	1.59	1.7	1.49	1.2		
PLACE1006360	1.1	5.11	1.79	2.46	2.74	2.62		

(1184)

特開2002-191363

2365

2366

PLACE1006364	4.51	8.06	7.29	7.37	9.19	5.75		
PLACE1006365	1.68	4.65	1.97	1.8	2.12	0.97		
PLACE1006368	1.53	3.11	2.57	3.01	3.04	4.27		
PLACE1006371	1.38	3.2	1.46	1.68	3.01	1.67		
PLACE1006373	2.21	5.21	5.75	7.83	8.02	7.56	*	+
PLACE1006382	0.9	4.67	2.81	3.3	1.92	2.95		
PLACE1006385	1.59	6.33	1.86	2.68	2.59	2.71		
PLACE1006391	1.19	5	1.95	1.96	2.79	1.63		
PLACE1006412	1.88	5.53	5.92	7.07	9.93	5.27		
PLACE1006414	0.63	3.42	0.95	1.22	1.87	1.6		
PLACE1006419	7.79	9.8	11.93	5.19	7.29	5.32	*	-
PLACE1006438	0.99	6.07	3.42	3.29	4.56	5.14		
PLACE1006443	2.05	5.01	5.12	5.01	5.31	6.44		
PLACE1006445	0.84	5.76	3.65	3.53	3.27	3.55		
PLACE1006447	1.34	5.81	3.28	2.95	3.26	3.96		
PLACE1006466	0.75	4.38	1.35	1.49	1.66	1.08		
PLACE1006469	0.67	4.66	2.31	1.65	2.26	1.67		
PLACE1006470	2.47	3.71	3.74	5.25	7.02	4.35		
PLACE1006472	24.4	23.44	52.17	26.23	28.52	9.36		
PLACE1006476	2.52	4.31	8.67	6.21	7.23	5.93		
PLACE1006482	1.64	3.35	4.43	4.25	4.67	4.98		
PLACE1006488	14.12	19.42	32.69	40.76	34.77	41.4	*	+
PLACE1006492	2.03	6.41	4.38	4.04	4.98	3.02		
PLACE1006506	1.78	6.67	4.04	4.41	5.71	4.17		
PLACE1006515	1.65	5.7	3.08	3.19	2.84	4.08		
PLACE1006516	1.1	7.32	7.05	4.89	5.69	7.28		
PLACE1006520	1.02	2.74	2.12	1.19	3	1.54		
PLACE1006521	2.4	3.54	6.38	6.49	6.86	5.08		
PLACE1006529	5.96	7.35	6.96	10.56	8.2	7.93		
PLACE1006531	1.01	4.31	3.33	1.84	2.05	2.43		
PLACE1006534	1.68	6.04	2.59	3.01	3.86	3.19		
PLACE1006540	2.68	9.7	7.77	8.71	11.21	4.46		
PLACE1006549	0.6	9.45	2.09	1.6	2.28	1.65		
PLACE1006550	1.76	8.82	4.07	2.77	2.94	4.14		
PLACE1006552	1.3	2.48	2.14	1.97	1.3	0.81		
PLACE1006557	2.38	4.01	3.79	2.84	2.51	2.45		
PLACE1006563	2.49	3.44	5.7	4.23	4.15	4.3		
PLACE1006579	1.53	7.5	4.82	4.88	5.38	5.78		
PLACE1006594	236.53	241.11	397.64	122.37	278.58	324.29		
PLACE1006598	0.72	8.53	2.4	1.53	1.58	2.07		
PLACE1006607	1.47	7.69	4.18	3.45	5.86	4.29		
PLACE1006610	9.46	13.73	38.26	27.65	32.76	22.64		
PLACE1006615	6.22	9.09	18.78	20.25	15.74	15.86		
PLACE1006617	0.91	1.54	2.66	1.87	2.49	2.09		
PLACE1006618	5.42	8.01	9.24	5.33	8.59	5.76		
PLACE1006626	1.53	4.11	1.3	2.47	2.78	1.16		
PLACE1006629	0.99	5.05	1.36	2.22	2.56	1.76		
PLACE1006637	1.29	6.54	3.97	3.77	4.23	4.87		
PLACE1006640	0.59	5.14	1.17	0.85	2.54	0.94		
PLACE1006644	1.66	4.46	2.12	2.79	2.49	2.39		

(1185)

特開 2002-191363

2367

2368

PLACE1006657	1.28	2.09	2.31	4.55	3.09	2.19		
PLACE1006673	2.29	4.73	10.34	11.06	10.89	6.45		
PLACE1006678	2.54	2.98	1.44	1.37	1.96	1.39		
PLACE1006682	3.5	5.93	2.58	15.44	20.96	23.99	**	+
PLACE1006684	1.12	4.8	1.81	1.64	2.54	1.65		
PLACE1006698	1.54	5.86	4.52	2.15	3.57	1.9		
PLACE1006704	1.81	5.41	2.71	2.93	2.92	2.97		
PLACE1006708	1.69	5.07	3.49	3.46	4.11	3.7		
PLACE1006711	14.21	16.18	29.77	24.34	26.25	22.42		
PLACE1006714	2.27	3.26	4.74	4.57	5.23	3.53		
PLACE1006716	1.51	2.75	3.7	6	7.05	3.99		
PLACE1006731	1.65	3.77	2.83	2.71	4	3.09		
PLACE1006754	0.43	3.94	1.73	1.8	1.81	0.99		
PLACE1006760	7.56	10.98	10.08	8.58	8.89	11.31		
PLACE1006779	1.44	4.12	2.88	3.19	3.79	2.97		
PLACE1006782	0.44	5.17	2.42	2.95	1.57	1.15		
PLACE1006783	9.34	11.46	18.65	157.98	223.05	66.46	*	+
PLACE1006786	3.31	4.08	6.07	5.9	6.24	3.34		
PLACE1006792	1.61	3.31	5.38	5.66	3.33	4.18		
PLACE1006795	0.89	2.43	0.74	0.81	1.27	1.01		
PLACE1006800	1.62	4.94	2.53	4.7	4.56	3.93		
PLACE1006805	3.94	7.79	5.5	10.83	9.8	8.79	*	+
PLACE1006809	3.55	5.7	5.94	9.58	10.61	8.97	**	+
PLACE1006815	1.7	7.57	4.1	5.12	5.23	5.8		
PLACE1006819	0.33	0.88	0.95	0.89	1.76	0.63		
PLACE1006820	2.35	2.01	4.91	4.84	6.72	4.18		
PLACE1006826	2.28	6.22	4.84	7.68	7.62	5.58		
PLACE1006829	3.76	5.51	6.54	9.49	8.66	8.69	*	+
PLACE1006853	1.2	4.21	1.97	2.25	2.93	2.88		
PLACE1006860	1	4.29	1.62	1.61	2.1	1		
PLACE1006867	5.65	9.36	11.34	7.04	8.33	7.63		
PLACE1006875	1.15	6.19	5.66	4.84	4.53	4.63		
PLACE1006878	1.59	2.84	3.09	2.99	3.22	2.39		
PLACE1006883	3.21	5.08	6.78	6.83	7.38	6.19		
PLACE1006898	1.67	4.23	3.67	3.54	4.77	4.59		
PLACE1006901	2.59	4.75	4.03	3.71	3.28	4.14		
PLACE1006904	0.91	3.59	2.7	3.26	2.92	2.04		
PLACE1006917	3.63	7.13	6.1	5.8	7.21	7.03		
PLACE1006932	0.54	5.85	1.29	0.92	1.34	1.19		
PLACE1006935	1.3	5.46	2.54	1.59	4.03	1.6		
PLACE1006956	0.92	2.55	3.4	2.55	2.41	2.09		
PLACE1006958	0.78	2.41	1.35	1.76	4.2	3.39		
PLACE1006959	4.97	8.48	9.98	11.46	9.58	13.62		
PLACE1006961	8.03	9.85	14.42	13.73	11.57	14.2		
PLACE1006962	2.97	7.44	6.56	5.04	7.26	6.22		
PLACE1006966	2.02	6.94	3.46	3.15	3.89	2.89		
PLACE1006979	0.95	4.44	2.03	1.46	2.64	1.77		
PLACE1006989	2.19	5.05	3.02	3.27	3.9	5.06		
PLACE1007001	4.98	6.79	10.71	4.03	7.43	7.38		
PLACE1007014	1.37	3.03	3.45	1.79	2.18	2.2		

(1186)

特開 2002-191363

2369

2370

PLACE1007021	0.74	3.03	2.11	0.75	2.2	1.73		
PLACE1007026	2.1	9.23	3.93	4.15	4.27	5.42		
PLACE1007028	4.12	8.5	10.56	7.89	8.34	9.35		
PLACE1007038	237.33	267.91	446.14	406.27	622.67	671.17		
PLACE1007040	1.55	3.14	2.85	1.57	3.31	2.45		
PLACE1007045	1.08	3.74	2.85	2.9	5.03	2.74		
PLACE1007048	147.06	149.67	259.53	121.61	211.26	109.43		
PLACE1007053	4.9	6.69	10	3.59	4.91	4.71		
PLACE1007068	7.56	10.33	62.76	39.52	45.9	36.69		
PLACE1007070	5.97	10.85	10.28	8.65	9.6	14.3		
PLACE1007076	8.22	14.4	14.19	16.53	23.62	24.67	*	+
PLACE1007077	2.65	6.45	4.01	5.2	5.28	5.43		
PLACE1007081	0.36	4.47	1.94	1.92	1.92	1.37		
PLACE1007082	1.23	4.66	4.95	4.32	4.5	3.99		
PLACE1007092	2.49	4.12	7.26	4.77	5.22	4.34		
PLACE1007096	0.72	2.19	0.74	1.35	1.63	0.97		
PLACE1007097	0.54	2.49	1.35	1.61	1.28	1.04		
PLACE1007099	1.58	4.66	2.56	2.77	3.64	3.72		
PLACE1007105	1.18	6.51	3.44	2.65	4.13	2.21		
PLACE1007108	3.55	13.02	7.41	5.03	6.87	5.75		
PLACE1007111	1.33	9.51	1.52	1.74	2.37	1.52		
PLACE1007112	1.23	7.26	1.79	2.09	3.12	2.36		
PLACE1007130	0.54	2.02	1.92	0.87	1.47	0.33		
PLACE1007132	1.46	3.32	4.63	3.58	3.38	2.88		
PLACE1007140	0.61	2.58	2.41	1.98	1.98	1.32		
PLACE1007143	2.79	6.32	4.62	4.9	5.34	5.33		
PLACE1007169	2.21	8.59	3.46	5.44	8.46	7.99		
PLACE1007178	0.82	8.66	2.48	3.28	6.28	4.1		
PLACE1007190	3.31	10.9	6.7	10.51	13.57	11.14		
PLACE1007201	0.81	5.82	1.41	1.72	3.04	2.51		
PLACE1007202	37.76	34.95	76.28	58.23	34.42	37.86		
PLACE1007226	2.01	2.39	2.73	1.89	3.14	2.29		
PLACE1007238	1.64	3.07	1.83	2.39	2.73	2.2		
PLACE1007239	1.81	3.68	2.99	1.76	2.72	2.44		
PLACE1007242	0.61	5.18	1.87	1.54	1.14	1.67		
PLACE1007243	2.21	7.36	2.29	2.24	3.27	3.31		
PLACE1007247	0.36	6.17	1.71	1.11	1.36	1.34		
PLACE1007257	1.67	5.33	3.34	3.3	5.27	4.25		
PLACE1007274	1.46	2.18	4.43	4.38	4.03	4.06		
PLACE1007276	0.93	2.02	1.1	2.13	2.1	1.74		
PLACE1007282	2.51	4.2	5.72	4.28	3.62	4.66		
PLACE1007286	2.97	4.8	7.85	10.14	12.47	8.79	*	+
PLACE1007296	10.55	19.45	24.46	31.43	17.57	27.05		
PLACE1007301	0.65	5.17	1.55	1.19	1.54	1.11		
PLACE1007314	3.11	6.61	8.64	7.98	8.96	10.24		
PLACE1007317	1.19	3.34	1.27	1.88	1.62	1.79		
PLACE1007329	0.89	0.73	1.78	2.38	2.35	2.09	*	+
PLACE1007338	3.96	6.47	9.58	11.59	8.93	2.32		
PLACE1007342	0.71	1.8	1.3	1.1	1.1	0.7		
PLACE1007345	1.72	4.57	2.54	2.72	3.6	3.29		

(1187)

特開 2002-191363

2371

2372

PLACE1007346	1.43	4.61	3.89	5.77	4.53	4.1		
PLACE1007359	0.74	4.55	2.16	2.59	2.44	3.53		
PLACE1007367	4.53	8.63	15.16	12.49	13.49	11.75		
PLACE1007375	0.36	3.24	2.02	1.75	2.56	1.59		
PLACE1007377	1.49	2.01	3.18	3.29	3.96	2.36		
PLACE1007386	1.55	1.75	1.47	2.37	1.68	1.36		
PLACE1007392	1.57	2.99	2.49	2.79	4.48	3.51		
PLACE1007402	2.41	5.66	3.08	1.52	2.91	1.8		
PLACE1007409	1.05	4.57	1.04	2.51	2.68	2.02		
PLACE1007416	3.45	6.97	6.5	7.05	9.14	6.52		
PLACE1007420	12.12	12.66	20.8	25.26	23.9	22.88	*	+
PLACE1007431	1.87	7.4	5.51	7.17	5.28	5.91		
PLACE1007450	0.79	1.22	2.65	3	2.99	2.39		
PLACE1007452	0.42	2.36	1.76	2.09	2.98	1.45		
PLACE1007454	23.74	28.02	76.56	59.97	75.95	46.61		
PLACE1007460	0.75	3.52	2.35	2.34	1.93	2.58		
PLACE1007478	0.41	3.07	1.33	1.35	2.18	1.92		
PLACE1007484	0.6	4.8	2.57	2.56	1.45	1.69		
PLACE1007488	0.4	6.24	1.64	1.74	2.61	1.46		
PLACE1007507	2.91	6.36	4.49	5.31	8.29	8.11		
PLACE1007511	0.53	1.29	1.06	1.06	1.29	0.42		
PLACE1007513	10.57	10.43	24.05	12.24	16.88	16.9		
PLACE1007524	1.55	3.33	3.53	3.96	4.72	2.96		
PLACE1007525	1.24	2.95	3.14	2.38	2.85	2.24		
PLACE1007537	8.6	9.68	49.88	43.78	63.66	40.1		
PLACE1007544	1.55	6.45	4.97	3.2	3.92	4.61		
PLACE1007547	1.36	5.03	4.15	2.37	2.84	2.4		
PLACE1007557	1.12	3.16	3.14	3.07	3.9	3.41		
PLACE1007560	9.38	8.86	12.57	11.03	9.62	17.59		
PLACE1007565	0.37	2.27	1	1	1.16	0.91		
PLACE1007580	1.06	3.71	3.06	10.8	11.15	13.74	**	+
PLACE1007583	0.76	3.88	1.78	2.51	2.37	1.09		
PLACE1007591	0.79	4.62	1.7	2.2	2.53	2.07		
PLACE1007598	1.13	6.98	3.86	2.71	3.46	4.71		
PLACE1007610	0.41	5.63	1.28	1.33	3.18	1.5		
PLACE1007618	1.57	1.91	2.01	1.75	2.2	2.41		
PLACE1007621	1.78	2.83	3.64	3.33	3.57	4.38		
PLACE1007626	23.99	25.61	32.78	30.53	30.94	13.53		
PLACE1007632	2.03	3.26	2.52	2.65	3.81	4.63		
PLACE1007635	1.61	4.62	6.42	2.8	4.19	3.37		
PLACE1007645	10.59	11.55	15.06	9.99	11.58	11.95		
PLACE1007649	1.7	5.88	3.47	2.78	4.95	3.13		
PLACE1007659	1.33	5.85	3.61	4.88	6.22	4.9		
PLACE1007669	2.01	2.1	3.74	2.97	4.63	4.4		
PLACE1007677	1.25	2.29	2.81	2.68	3.07	2.91		
PLACE1007688	3.4	5.69	5.43	1.98	4.53	4.98		
PLACE1007690	1.4	4.03	2.12	3.74	3.37	4.61		
PLACE1007697	0.69	7.13	1.37	1.84	2.56	1.8		
PLACE1007702	2.03	7.08	5.7	4.03	3.91	4.08		
PLACE1007705	1.38	3.93	1.59	1.74	4.45	2.75		

(1188)

特開2002-191363

2373

2374

PLACE1007706	3.11	6.08	4.69	5.25	8.84	7.49	
PLACE1007725	3.41	4.69	6.65	5.1	3.29	5.39	
PLACE1007729	0.98	2.65	1.8	2.7	3.11	1.99	
PLACE1007730	1.25	4.29	3.07	3.66	3.75	4.32	
PLACE1007737	1.43	4.79	3.39	3.79	4.7	4.17	
PLACE1007743	1.38	4.26	2.29	2.3	2.83	2.03	
PLACE1007746	6.56	9.02	10.42	9.65	13.29	12.97	
PLACE1007753	0.53	4.48	1.71	1.35	2.86	1.94	
PLACE1007769	1.31	4.31	3.5	3.27	4.51	4.58	
PLACE1007780	5.77	4.63	7.11	6.51	3.75	2.17	
PLACE1007791	1.82	3.29	3.38	3.16	3.69	2.87	
PLACE1007807	0.67	2.79	1.72	2.33	1.76	1.29	
PLACE1007810	0.39	4.45	2.63	4.11	4.08	3.27	
PLACE1007814	3.57	5.98	5.04	4.2	4.62	6.3	
PLACE1007828	2.01	7.64	3.34	2.69	4.64	3.44	
PLACE1007829	1.32	6.9	2.88	2.87	4.87	3.06	
PLACE1007841	1.64	7.26	1.87	2.25	3.14	3.39	
PLACE1007842	1.1	3.32	2.44	2.09	3.96	1.39	
PLACE1007843	1.2	1.92	1.43	2.13	1.48	1.86	
PLACE1007845	1.76	3	4.11	3.45	3.42	2.36	
PLACE1007846	0.99	3.26	1.64	2.02	2.73	1.5	
PLACE1007848	1.09	3.51	2.23	2.39	2.62	2.25	
PLACE1007852	2.26	7.88	3.82	2.94	4.61	3.24	
PLACE1007858	3.65	11.57	5.81	61.71	80.46	57.09	** +
PLACE1007866	19.42	25.98	40.48	43	80.39	56.73	
PLACE1007871	8.1	7.9	15.45	16.17	12.35	11.08	
PLACE1007877	1.09	2.09	1.45	1.4	2.39	1.53	
PLACE1007878	5.98	9.75	14.61	13.65	7.49	8.9	
PLACE1007881	0.43	2.66	1.34	1.59	1.94	1.93	
PLACE1007885	4.35	7.85	6.76	5.57	6.53	7.01	
PLACE1007897	0.27	6.51	1.85	1.72	1.53	1.41	
PLACE1007908	3.14	12.29	5.73	5.96	7.9	8.24	
PLACE1007922	6.08	11.75	8.75	5.24	7.15	4.54	
PLACE1007946	1.07	2.03	1.86	2.71	2.28	1.94	
PLACE1007950	6.98	7.6	18.21	16.17	19.34	12.63	
PLACE1007954	-0.03	2.45	1.15	2.46	1.57	1.51	
PLACE1007955	0.92	4.01	2.17	2.05	2.52	3.05	
PLACE1007956	0.6	3.61	2.91	2.35	2.1	2.22	
PLACE1007958	0.75	6.31	1.34	0.79	1.21	0.8	
PLACE1007965	0.64	5.88	3.25	3.38	2.91	2.17	
PLACE1007969	1.09	6.37	3.06	2.35	3.29	2.21	
PLACE1007971	2.73	4.17	5.21	6.1	4.41	5.92	
PLACE1007990	1.95	2.33	2.31	3.22	3.09	1.88	
PLACE1008000	0.32	2.16	1.98	1.85	1.27	0.66	
PLACE1008002	0.99	3.38	1.7	1.81	2.04	0.51	
PLACE1008037	0.57	4.19	1.7	4.59	2.86	2.02	
PLACE1008044	1.42	5.81	2.46	4.18	4.93	4.16	
PLACE1008045	0.4	4.07	1.54	1.75	2	1.65	
PLACE1008080	2.05	6.08	3.22	4.23	4.03	4.78	
PLACE1008092	1.56	1.56	1.48	1.48	2.98	1.86	

(1189)

特開 2002-191363

2375

2376

PLACE1008095	0.59	2.14	1.48	2.38	2.73	1.23		
PLACE1008105	0.95	1.76	1.71	2.24	2.71	0.74		
PLACE1008107	0.27	2.33	0.7	1.72	1.44	1.68		
PLACE1008111	1.73	5.01	2.12	4.57	5.4	4.04		
PLACE1008113	5.88	9.24	12.48	16.57	20.29	19.24	*	+
PLACE1008122	1.22	5.54	2.55	1.61	1.57	1.5		
PLACE1008129	1.5	5.64	2.8	2.43	5.36	2.91		
PLACE1008132	5.51	4.47	8.34	6.61	11.2	7.63		
PLACE1008137	0.96	1.82	1.02	2.12	3.88	0.8		
PLACE1008174	0.77	3.16	2.43	5.12	4.39	2.46		
PLACE1008177	1.62	4.87	3.09	3.79	3.26	3.77		
PLACE1008181	1.76	3.87	1.6	2.06	2.43	1.43		
PLACE1008195	2.66	6.08	2.97	4.34	4.14	3.9		
PLACE1008198	1.06	5.56	2.52	2.55	3.31	1.54		
PLACE1008201	1.22	4.45	3.58	5.92	7.69	5.64	*	+
PLACE1008209	2.35	2.29	4.46	2.51	5.31	4.44		
PLACE1008226	1.8	2.35	5.25	4.72	5.68	7.08		
PLACE1008227	0.77	2.67	3.02	2.38	4.6	3.54		
PLACE1008231	1.26	3.85	1.85	1.05	1.83	0.78		
PLACE1008238	1.22	3.21	2.9	2.47	2	1.72		
PLACE1008244	1.01	4.69	1.56	1.68	3.2	1.56		
PLACE1008249	0.8	4.94	2.55	1.22	2.17	2.01		
PLACE1008266	11.31	18.61	43.04	60.04	82.48	59.04	*	+
PLACE1008273	1.47	3.95	3.81	3.53	3.8	4.47		
PLACE1008275	1.59	3.67	2.17	2.62	2.57	2.34		
PLACE1008280	0.85	2.6	1.84	2.42	2.48	2.36		
PLACE1008282	4.71	8.19	6.89	7.27	9.02	6.38		
PLACE1008297	2.32	4.7	3.36	2.89	3.42	3.21		
PLACE1008303	1.65	6.68	1.24	4.12	3.83	2.65		
PLACE1008309	0.43	6.52	0.82	1.77	1.5	1.29		
PLACE1008315	5.3	5.93	8.61	4.92	4.79	9.83		
PLACE1008329	0.47	2.23	2.06	2.32	2.8	2.49		
PLACE1008330	0.72	4.06	3.16	2.48	3.36	2.96		
PLACE1008331	0.84	5.01	2.1	4.5	2.17	2.91		
PLACE1008351	4.34	8.66	7.41	7.91	7.31	7.1		
PLACE1008356	1.56	8.23	1.93	2.86	4.16	3.35		
PLACE1008359	1.57	4.11	2.89	2	2.97	2.94		
PLACE1008368	2.27	6.38	7.43	5.72	7.33	6.95		
PLACE1008369	0.57	2.46	1.45	1.12	1.59	1.68		
PLACE1008392	0.8	3.09	2.54	2.44	3.22	3.24		
PLACE1008394	2.08	4.84	3.75	3.98	5.03	4.76		
PLACE1008398	5.32	9.36	11.44	11.36	11.3	12.33		
PLACE1008401	1.19	7.06	3.21	2.82	3.43	3.33		
PLACE1008402	3.21	6.45	7.2	7.23	10.15	9.26		
PLACE1008405	10.3	10.95	18.42	17.17	18.82	20.4		
PLACE1008409	1.88	5.19	5.69	4.97	5.41	5.65		
PLACE1008420	1.4	1.87	1.96	2.67	2.69	2.27	*	+
PLACE1008424	0.88	2.69	2.54	1.69	2.34	1.71		
PLACE1008426	0.98	2.58	1.58	1.7	2.66	2.32		
PLACE1008429	0.92	3.17	2.14	1.91	3.4	1.84		

(1190)

特開2002-191363

2377

2378

PLACE1008430	1.63	4.85	3.04	2.93	3.52	3	
PLACE1008437	0.87	3.64	3.01	2.83	1.82	1.57	
PLACE1008453	1.16	4.8	1.02	1.64	2.06	1.17	
PLACE1008454	2.14	6.46	9.23	5.46	9.02	5.92	
PLACE1008455	2.06	4.33	7.2	5.26	6.68	4.87	
PLACE1008457	0.51	2.6	2.01	2.28	2.43	2.47	
PLACE1008465	0.49	2.41	1.72	1.56	2.13	0.48	
PLACE1008469	2.42	4.36	5.32	5.16	4.75	7.1	
PLACE1008488	0.81	5.48	1.97	2.44	2.03	1.8	
PLACE1008519	1.48	10.85	6.17	4.41	4.99	4.51	
PLACE1008524	1.04	11.09	1.72	2	3.22	2.02	
PLACE1008531	0.64	8.37	1.92	1.33	2.05	1.72	
PLACE1008532	2.12	3	5.51	5.66	4.72	4.19	
PLACE1008533	2.01	4	4.07	5.53	5.18	3.77	
PLACE1008542	1.61	2.36	0.96	2.05	2.1	1.72	
PLACE1008549	0.96	3.06	0.67	1.45	2.1	1.53	
PLACE1008560	1.18	4.23	2.28	2.29	3.93	3.47	
PLACE1008567	0.87	7.26	1.85	2.33	3.67	2.38	
PLACE1008568	2.37	10.67	5.49	2.97	7.47	4.21	
PLACE1008569	3.94	10.32	6.74	6.1	8.6	7.6	
PLACE1008584	0.88	1.4	1.58	2.86	3.38	1.31	
PLACE1008585	4.96	4.8	7.56	11.08	4.84	3.57	
PLACE1008603	5.9	7.25	31	30.55	43.67	29.76	
PLACE1008621	0.55	2.28	0.95	0.72	1.89	1.16	
PLACE1008625	0.64	4.01	0.9	1.18	1.41	2.03	
PLACE1008626	0.55	6.06	0.9	1.03	0.83	0.95	
PLACE1008627	0.46	8.32	1.86	1.87	3.34	2.7	
PLACE1008629	3.22	9.18	5.84	5.44	6.75	4.41	
PLACE1008630	1.68	3.39	4.23	4.21	3.01	3.51	
PLACE1008643	1.31	0.93	1.98	1.72	2.34	1.94	
PLACE1008650	0.25	3.05	1.62	2.23	1.63	1.24	
PLACE1008657	1.17	2.39	2.51	2.34	4.04	2.91	
PLACE1008664	0.91	5.93	2.37	2.91	2.51	1.13	
PLACE1008693	0.97	4.93	3.09	2.53	3.81	2.2	
PLACE1008696	0.88	3.84	2.21	2.26	2.11	1.47	
PLACE1008715	1.05	4.71	2.11	1.34	2.65	2.65	
PLACE1008716	2.48	3.94	4.19	5.75	6.9	7.07	* +
PLACE1008722	3.85	4.34	7.37	7.64	7.19	3.45	
PLACE1008738	5.17	9.13	12.7	9.49	5.83	4.8	
PLACE1008742	6.57	6.87	14.66	14.94	15.06	12.41	
PLACE1008744	3.52	6.98	5.61	5.83	4.55	2.74	
PLACE1008748	0.63	4.39	2.75	2.44	1.67	1.61	
PLACE1008757	0.99	4.74	4.51	2.77	5.74	2.17	
PLACE1008766	2.66	6.75	3.77	3.51	6.47	4.06	
PLACE1008785	1.39	1.68	2.6	3.26	3.8	3.89	* +
PLACE1008790	1.57	1.8	2.29	3.5	5.39	2.96	
PLACE1008798	1.71	3.82	4.45	6	5.93	3.32	
PLACE1008807	1.34	3.95	1.61	2.54	2.62	1.8	
PLACE1008808	1.6	4.53	3.01	4.24	3.69	5.04	
PLACE1008813	1.38	4.85	1.97	1.9	1.95	2.3	

(1191)

特開 2002-191363

2379

2380

PLACE1008836	1.34	5.81	3.68	3.89	6.17	4.1
PLACE1008851	1.21	6.65	3.94	4.85	10.04	4.54
PLACE1008854	0.56	0.48	1.14	1.16	1.41	1.6
PLACE1008864	1.98	1.92	2.96	2.73	2.65	2.49
PLACE1008867	1.2	6.57	6.22	6.43	5.71	4.82
PLACE1008876	11.7	16.5	27.29	26.74	20.94	23.24
PLACE1008887	1.37	4.31	1.44	3.26	2.07	3.28
PLACE1008902	1.33	5.62	2.93	3.17	4.58	2.06
PLACE1008911	4.04	8.56	10.48	11.31	13.99	15.66
PLACE1008917	0.6	4.53	2.72	1.7	2.9	1.71
PLACE1008920	0.75	0.77	0.87	0.61	1.58	1.44
PLACE1008925	0.25	0.9	1.04	0.94	1.91	0.84
PLACE1008930	4.12	7.32	9.83	5.11	10.36	7.17
PLACE1008934	0.9	3.42	2.9	2.89	2.28	1.7
PLACE1008941	1.57	4.14	2.8	2.06	2.59	4.05
PLACE1008947	2.3	5.41	5.51	3.96	5.84	5.16
PLACE1008984	1.26	6.31	3.25	3.1	3.93	3.19
PLACE1008985	0.94	2.75	2.74	2.84	2.43	2.7
PLACE1008994	0.27	1.72	0.65	1.11	0.78	0.68
PLACE1009020	0.46	3.49	2.42	2.49	3.1	2.16
PLACE1009027	0.89	2.7	1.59	2.24	1.75	2.09
PLACE1009039	-0.06	3.31	3.42	2.39	1.59	1.49
PLACE1009045	1.53	6.33	6.05	23.13	20.76	22.2
PLACE1009048	0.41	5.97	2.3	0.61	1.04	0.54
PLACE1009050	0.97	4.9	1.68	1.07	1.47	1.55
PLACE1009060	5.61	8.4	9.51	10.74	8.55	11.96
PLACE1009067	1.14	2.8	2.03	1.6	2.34	3.4
PLACE1009071	1.44	4.05	3.9	3.79	7.24	9.82
PLACE1009090	1.27	6.46	2.35	3.11	4.73	2.86
PLACE1009091	5.58	10.22	38.11	38.77	49.35	36.29
PLACE1009094	0.26	5.68	1.88	1.67	5.04	1.71
PLACE1009099	1	5.52	3.47	3.49	3.36	3.84
PLACE1009110	1.59	5.82	1.16	1.68	4.3	4.39
PLACE1009111	1.88	5.24	2.65	3.95	3.77	2.88
PLACE1009113	2.24	3.52	3.62	6.14	4.87	7.29
PLACE1009130	4.46	6.8	7.84	6.68	9.36	10.47
PLACE1009150	0.88	3.54	1.95	3.23	3.3	3.01
PLACE1009155	1.11	5.06	2.98	4.46	4.43	3.87
PLACE1009158	1.06	5.77	1.95	1.77	2.35	2.88
PLACE1009166	0.76	4.8	1.53	1.59	2.16	1.3
PLACE1009172	1.43	3.96	2.45	2.26	5.85	2.61
PLACE1009174	1.13	3.45	2.42	1.67	3.02	2.38
PLACE1009183	1.62	3.54	4.47	4.1	6.33	8.06
PLACE1009186	1.04	5.07	2.3	2.46	2.86	2.91
PLACE1009190	0.75	2.32	1.44	1.53	1.9	1.65
PLACE1009196	0.81	4.01	2.73	2.24	2.38	1.99
PLACE1009200	1.01	4.44	2.94	2.84	4.39	2.91
PLACE1009217	2.55	4.91	3.43	4.46	7.29	7.23
PLACE1009230	1.9	5.55	6.63	5.63	9.16	9.46
PLACE1009236	4.97	7.07	12.6	8.21	10.79	7.13

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(1192)

特開 2002-191363

2381

2382

PLACE1009246	11.71	11.96	24.75	14.59	16.36	9.05		
PLACE1009265	6.95	7.82	14.01	15.61	5.19	12.17		
PLACE1009279	0.67	2.07	2.46	1.93	2.54	1.63		
PLACE1009298	5.54	9.92	9.52	10.21	11.25	17.55		
PLACE1009308	1.13	6.82	2.04	2.48	2.48	2.34		
PLACE1009319	2.04	9.25	3.15	2.92	3.54	2.5		
PLACE1009328	1.04	5.78	1.81	2.98	3.39	2.17		
PLACE1009335	1.38	6.55	4.72	2.24	3.21	3.01		
PLACE1009338	2.56	4.14	5.1	3.24	4.3	1.57		
PLACE1009344	0.73	2.45	1.08	1.31	1.55	0.84		
PLACE1009355	5.41	7.37	9.95	13.44	10.76	13.55	*	+
PLACE1009368	1.3	2.56	2.41	2.43	2.32	2.19		
PLACE1009375	1.21	6.41	3.05	3.04	4.46	2.53		
PLACE1009388	1.18	8.68	3.01	3.46	4.53	2.72		
PLACE1009398	1.19	9.2	3.74	3.17	4.28	3.96		
PLACE1009404	2.78	9.18	4.51	5.33	6.73	6.94		
PLACE1009410	1.27	2.35	2.33	2.51	2.31	1.44		
PLACE1009417	0.95	2.25	4.34	2.55	3.08	1.71		
PLACE1009424	1.88	3.61	3.18	2.85	3.24	3.93		
PLACE1009434	0.84	3.94	2.91	1.29	1.82	2.19		
PLACE1009443	1.21	7.2	2.55	2.42	3.43	3.17		
PLACE1009444	1.33	7.71	4.05	2.51	3.17	3.79		
PLACE1009459	0.23	7.99	1.55	1.71	1.83	0.86		
PLACE1009460	1.75	6.84	3.26	5.15	4.31	4.08		
PLACE1009468	0.99	2.83	3.42	4.43	4.42	2.97		
PLACE1009476	0.21	1.21	0.73	1.05	0.67	1.33		
PLACE1009477	1.35	3.13	2.67	3.06	2.35	2.2		
PLACE1009493	0.87	3.35	0.94	1	1.87	1.41		
PLACE1009502	0.76	4.64	2.13	1.19	1.89	1.66		
PLACE1009524	1.32	4.22	1.63	0.94	2.14	1.6		
PLACE1009527	0.95	4.51	2.11	1.64	2.55	1.28		
PLACE1009531	20.82	28.24	41.52	46.25	43.25	49.96		
PLACE1009535	1.1	1.56	2.68	2.42	2.15	1.11		
PLACE1009539	2.15	3.41	4.18	3.88	2.65	2.57		
PLACE1009540	5.89	8	11.66	14.8	4.47	3.84		
PLACE1009542	1.11	3.37	1.42	1.51	2.06	1.44		
PLACE1009546	0.62	5.27	0.97	2.24	1.64	1.25		
PLACE1009556	0.35	4.46	3.46	3.36	2.86	3.16		
PLACE1009569	0.05	3.93	2.46	3.34	2.7	3.11		
PLACE1009571	1.67	4.27	2.52	3.04	2.85	2.67		
PLACE1009573	3.81	2.97	6.73	6.92	8.12	6.49		
PLACE1009576	1.92	2.51	4.3	2.73	3.66	2.08		
PLACE1009580	1.42	1.81	1.74	2.73	3.47	2.33	*	+
PLACE1009581	0.89	4.25	2.03	2.91	4.38	2.74		
PLACE1009587	0.96	4.91	2.29	2.43	3.2	1.99		
PLACE1009593	2.71	6.73	4.37	4.94	6.85	5.03		
PLACE1009595	1.81	5.44	2.66	2.67	5	2.79		
PLACE1009596	1.57	6.83	2.6	3.44	3.97	2.7		
PLACE1009600	3.03	4.27	4.48	5.48	9.14	4.42		
PLACE1009604	2.32	4.64	5.02	4.22	6.11	3.23		

(1193)

特開2002-191363

2383

2384

PLACE1009607	1.29	2.48	3.18	3.19	4.17	3.18
PLACE1009613	1.94	5.23	2.94	2.65	3.08	2.23
PLACE1009621	1.66	6.72	3.32	8.21	8.67	8.06
PLACE1009622	1.78	5.93	3.78	3.9	4.1	3.9
PLACE1009624	1.16	5.77	3.42	3.2	3.65	3.5
PLACE1009637	2	6.88	3.36	3.07	4.59	3.91
PLACE1009639	1.94	1.76	4.15	3.44	3.67	4.99
PLACE1009654	20.88	17.13	34.95	14.94	24.53	20.64
PLACE1009659	2.77	6.78	7.45	6.38	8.38	6.55
PLACE1009665	1.04	4.21	1.93	1.19	2.72	1.93
PLACE1009669	7.73	9.64	14.54	9.85	16.89	8.82
PLACE1009670	1.76	5.36	2.54	2.77	4.47	4.01
PLACE1009708	2.1	5.57	5.09	3.64	6.54	5.84
PLACE1009721	1.34	4.28	3.56	5.78	5.81	3.01
PLACE1009731	1.36	3.59	3	3.58	6.53	5
PLACE1009735	1.94	3.94	3.21	5.16	7.52	4.78
PLACE1009737	1.89	4.29	2.95	4.83	5.61	5.47
PLACE1009741	1.3	4.32	3.45	2.09	5.03	3.07
PLACE1009752	1.34	5.64	2.65	2.3	3.33	1.68
PLACE1009763	3.95	9.73	6.82	7.13	7.44	8.39
PLACE1009766	1.46	6.98	3.32	3.07	5.19	3.75
PLACE1009772	0.48	5.19	0.6	1.01	2.46	0.89
PLACE1009782	0.91	2.39	2.03	2.88	2.91	3.74
PLACE1009794	2.58	4.45	5.11	3.54	3.66	5.03
PLACE1009798	1.59	5.37	4	6.26	5.57	5.67
PLACE1009845	1.05	6.02	2.92	2.79	3.39	3.92
PLACE1009849	0.96	6.61	2.35	1.79	3.41	2.59
PLACE1009857	0.79	4.86	1.45	1.19	1.27	1.56
PLACE1009861	1.43	4.67	3.87	4.1	3.47	3.11
PLACE1009872	53.53	52.43	88.5	74.95	49.47	81.73
PLACE1009877	5.45	7.59	12.08	10.03	10.3	12.32
PLACE1009879	0.82	3.28	1.59	1.55	2.99	1.34
PLACE1009886	0.68	4.04	1.53	1.62	3.04	1.72
PLACE1009888	1.03	7.4	3.23	5.34	5.84	7.94
PLACE1009908	1.56	7.63	8.64	3.37	6.71	5.9
PLACE1009919	4.5	7.53	8.26	5.84	10.72	10.15
PLACE1009921	0.96	3.94	3.32	1.63	4.28	2.47
PLACE1009923	3.82	5.56	6.85	6.32	8.13	5.57
PLACE1009924	3.01	2.49	4.53	4.43	4.31	1.04
PLACE1009925	0.61	2.77	1.84	2.51	2.2	2.5
PLACE1009931	2.78	5.21	9	8.71	6.93	8.09
PLACE1009935	0.74	3.71	2.1	1.19	1.08	1.5
PLACE1009947	0.47	3.83	1.64	1.51	2.46	2.03
PLACE1009961	0.43	4.08	1.39	1.69	2.18	1.9
PLACE1009971	0.92	4.9	1.98	1.35	1.74	1.45
PLACE1009982	40.34	48.71	89.8	53.8	57.87	66.96
PLACE1009992	0.94	1.9	2.59	1.47	2.52	0.68
PLACE1009995	6.47	10.83	15.72	7.79	9.03	11.23
PLACE1009997	0.55	3.7	3.03	2.76	3.2	2.64
PLACE1010002	1.4	4.14	2.82	2.89	3.04	3.46

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(1194)

特開2002-191363

2385

2386

PLACE1010011	2.09	8.13	3.85	4.4	5.21	4.68		
PLACE1010013	0.18	12.85	1.74	0.92	1.81	0.68		
PLACE1010021	3.18	11.98	5.42	4.12	4.13	6.06		
PLACE1010023	2.15	8.45	5.16	5.62	6.52	6.14		
PLACE1010031	4.6	4.35	7.23	6.79	4.91	1.82		
PLACE1010039	1.17	3.3	1.45	1.28	1.23	1.19		
PLACE1010045	1.1	2.83	3.66	2.52	3.55	2.64		
PLACE1010053	1.42	3.56	1.65	2.21	2.76	2.37		
PLACE1010060	1.63	6.1	4.13	4.11	4.6	4.05		
PLACE1010069	0.41	7.96	2.32	1.48	2.91	1.3		
PLACE1010070	0.92	8.04	1.5	0.45	1.78	1.09		
PLACE1010074	5.25	11.67	11.8	12.32	9.51	11.22		
PLACE1010076	12.75	11.95	29.01	19.58	15.88	16.82		
PLACE1010078	2.96	2.42	4.36	4.64	4.39	3.85		
PLACE1010081	2.74	4.1	3.7	6.85	7.81	4.59	*	+
PLACE1010083	0.69	2.53	1.51	1.22	1.86	2.26		
PLACE1010089	1.86	4.89	3.35	3.45	3.47	3.48		
PLACE1010096	2.17	7.73	2.91	3.43	4.19	3.2		
PLACE1010102	3.89	10.9	5.33	7.1	9.64	7.57		
PLACE1010105	2.98	7.46	4.93	7.31	9.04	9.82		
PLACE1010106	2.46	2.95	4.48	5.47	4.88	5.8	*	+
PLACE1010130	0.53	1.79	1.17	2.27	2.55	1.23		
PLACE1010132	2.49	4.65	5.3	5.07	4.39	4.19		
PLACE1010134	0.8	3.32	1.15	1.97	2.31	1.67		
PLACE1010139	6.67	10.51	12.98	14.99	16.1	14.58		
PLACE1010148	0.96	5.07	1.62	1.48	1.9	1.97		
PLACE1010152	3.11	5.68	5.16	6.33	6.64	5.76		
PLACE1010155	3.8	6.17	6.52	16.85	20.56	20.32	**	+
PLACE1010156	13.71	15.43	32.21	132.45	85.59	134.99	**	+
PLACE1010161	1.9	2.81	5.05	3.29	2.92	1.97		
PLACE1010181	0.73	2.22	1.51	2.58	1.99	2.53		
PLACE1010194	0.64	3.35	1.03	2.26	2.14	1.64		
PLACE1010202	0.4	4.14	1.2	2.91	1.65	2.16		
PLACE1010231	1.1	3.78	2.39	1.31	2.99	2.73		
PLACE1010235	1.26	4.24	1.94	2.68	2.42	3.16		
PLACE1010237	1.01	3.4	2.1	1.14	1.97	0.87		
PLACE1010251	0.59	0.98	1.95	2.57	3.18	1.62		
PLACE1010261	0.97	2.63	2.07	2.69	1.69	1.55		
PLACE1010270	0.76	2.7	1.3	1.39	2.33	1.42		
PLACE1010273	0.97	3.27	0.46	1.48	2.25	1.5		
PLACE1010274	6.28	9.23	9.66	10.49	12.18	14.28		
PLACE1010277	6.03	10.14	12.68	14.6	16.06	15.84	*	+
PLACE1010293	1.8	5.68	3.55	3.65	3.37	3.96		
PLACE1010297	5.17	11.37	21	24.84	32.59	22.06		
PLACE1010300	4.18	4.78	8.22	8	9.95	6.87		
PLACE1010310	16.52	14.75	49.45	70.74	71	77.01	*	+
PLACE1010321	2.03	4.92	2.46	3.37	4.99	2.73		
PLACE1010324	0.88	3.49	1.56	1.12	1.54	1.2		
PLACE1010329	0.73	4.64	1.95	2.56	3.37	1.96		
PLACE1010330	3.78	9.09	7.29	2.42	10.45	7.67		

(1195)

特開2002-191363

2387

2388

PLACE1010335	6.43	11.15	7.43	13.15	17.89	19.1	*	+
PLACE1010341	0.19	4.81	1.24	1.07	3.13	1.54		
PLACE1010342	0.77	0.9	0.75	0.48	2.12	0.8		
PLACE1010346	1.47	1.73	3.61	2.96	4.47	1.71		
PLACE1010362	1.31	2.69	2.22	2.18	3.49	3.31		
PLACE1010364	0.78	2.56	1.65	1.19	2.32	1.49		
PLACE1010368	1.66	5.44	3.51	3.41	3.87	4.48		
PLACE1010373	9.05	10.48	16.82	12.13	15.45	12.28		
PLACE1010383	1.91	5.52	5.13	5.58	6.39	4.9		
PLACE1010385	0.3	3.01	1.07	0.04	0.6	0.9		
PLACE1010389	6.28	7.98	13.24	22.3	13.64	22.94	*	+
PLACE1010401	0.73	2.72	1.32	1.99	2.87	2.21		
PLACE1010410	3.15	4.83	6.71	4.78	7.55	7.32		
PLACE1010418	1.88	4.73	4.2	4.71	5.14	4.66		
PLACE1010425	0.93	4.78	1.43	1.78	1.96	2.15		
PLACE1010443	6.98	13.83	51.39	36.22	63.67	48.93		
PLACE1010445	0.95	5.02	0.68	2.69	3.81	2.44		
PLACE1010481	1.19	2.06	2.46	1.75	2	1.85		
PLACE1010482	28.99	29.39	53.06	31.75	19.63	40.44		
PLACE1010491	3.36	6.4	5.38	4.96	5.98	3.26		
PLACE1010492	4	4.75	5.95	5.57	8.15	7.77		
PLACE1010509	0.8	4.32	3.15	3.09	3.01	2.99		
PLACE1010518	3.33	6.72	6.58	7.3	8.25	8.24		
PLACE1010522	2.3	5.96	4.35	1.87	3.52	2.64		
PLACE1010529	1.8	6.5	3.84	5.43	8.44	7.2		
PLACE1010547	0.79	1.57	1.76	1.51	1.93	2.45		
PLACE1010560	0.63	2.51	2.06	3.31	2.2	2.51		
PLACE1010562	0.74	2.68	1.65	1.64	2.1	1.63		
PLACE1010579	1.11	7.13	2.63	3.45	4.67	3.26		
PLACE1010580	1.35	9.12	3.79	3.27	2.49	4.52		
PLACE1010599	3.56	6.07	7.94	8.32	9.26	8.13		
PLACE1010606	1.17	4.42	1.64	3.39	4.17	3.51		
PLACE1010616	1.84	3.72	5.49	4.09	8.09	6.81		
PLACE1010622	2.1	3.43	3.91	5.07	4.74	4.69	*	+
PLACE1010624	1.43	3.35	3.98	4.21	6.17	5.22		
PLACE1010628	1.37	3.97	2.83	2.14	3.63	4.22		
PLACE1010629	1.08	4.64	2.24	3.01	3.3	3.5		
PLACE1010630	1.64	4.77	3.69	4.22	4.78	5.29		
PLACE1010631	0.5	5.35	2.64	1.55	1.66	2.67		
PLACE1010651	14.24	15.75	24.44	37.62	40.09	52.12	*	+
PLACE1010661	1.62	4.09	2.28	3.56	6.43	3.22		
PLACE1010662	1.32	2.48	1.6	2.98	2.3	1.61		
PLACE1010668	12.87	15.91	27.82	37.63	30.53	28.75		
PLACE1010702	1.46	2.34	4.24	3.59	3.6	3.2		
PLACE1010709	79.16	78.33	115.91	107.07	96.3	133.25		
PLACE1010713	7	10.81	14.7	9.14	8.16	15.14		
PLACE1010714	0.82	7.41	1.58	1.75	2.04	1.47		
PLACE1010716	0.71	6.19	4.31	2.08	2.3	1.93		
PLACE1010717	0.9	6.49	2.13	2.17	3.9	2.61		
PLACE1010720	14.03	17.05	53.79	46.72	50.7	41.49		

(1196)

特開2002-191363

2389

2390

PLACE1010739	0.9	1.2	1.11	1.73	1.21	1.93		
PLACE1010743	1.09	2.3	1.99	2.63	2.05	2.21		
PLACE1010752	0.87	2.92	1.85	1.76	2.05	1.53		
PLACE1010761	3.6	8.83	13.51	12.07	16.4	17.08		
PLACE1010771	1.41	6.89	5.03	6.13	10.3	5.42		
PLACE1010784	0.9	9.66	1.97	2.07	1.72	1.34		
PLACE1010786	1.21	8.77	2.83	2.91	3.68	2.21		
PLACE1010789	0.6	1.16	1.52	1.8	1.89	1.17		
PLACE1010800	2.18	2.86	3.25	3.95	3.24	2.93		
PLACE1010802	2.97	4.63	5.31	5.72	4.27	3.3		
PLACE1010811	0.89	2.19	1.96	1.83	1.75	2.18		
PLACE1010813	8.89	13.3	55.85	48.82	72.26	46.7		
PLACE1010827	1.54	6.43	3.94	4.3	5.52	4.81		
PLACE1010833	0.93	8.13	2.63	2.68	3.64	2.09		
PLACE1010839	1.57	6.22	3.21	4.22	6.72	4.13		
PLACE1010856	7.58	8.94	12.34	8.02	6.53	8.59		
PLACE1010857	3.41	3.81	7.63	8.24	5.98	4.56		
PLACE1010870	1.3	2.24	2.05	1.62	2.38	1.94		
PLACE1010877	1.67	4.66	2.69	2.77	3.92	2.62		
PLACE1010882	0.49	4.8	0.99	1.74	1.27	0.51		
PLACE1010891	1.1	7.73	1.12	1.85	1.73	0.95		
PLACE1010896	1.19	5.29	3.18	3.98	3.05	3.47		
PLACE1010900	7.41	13.29	27.9	23.88	20.99	18.85		
PLACE1010916	1.55	1.18	2.06	1.89	1.9	2.2		
PLACE1010917	-0.04	0.82	0.36	1.05	2.25	0.56		
PLACE1010924	1.15	2.31	1.55	1.11	1.88	0.92		
PLACE1010925	2.76	5.36	2.17	4.92	6.32	4.16		
PLACE1010926	1.8	5.73	4.31	5.37	4.35	4.45		
PLACE1010942	1.7	6.25	5.63	5.53	7.88	7.69		
PLACE1010943	7.38	10.43	17.12	24.62	29.5	31.96	**	+
PLACE1010944	4.33	7.39	9.3	13.11	11.44	15.58	*	+
PLACE1010947	1.43	0.9	2.41	2.57	2.8	2.25		
PLACE1010954	3.56	2.92	7.4	7.92	8.55	7.64		
PLACE1010960	2.06	3.44	6.07	4.23	7.89	2.6		
PLACE1010965	2.33	3.81	3.54	6.09	4.92	5.03	*	+
PLACE1010968	1.55	4.69	1.38	4.95	6.51	3.68		
PLACE1010978	3.63	6.12	7.05	9	8.94	6.45		
PLACE1010982	2.23	5.77	5.6	4.74	5.66	4.88		
PLACE1010990	0.88	5.4	2.04	3.11	3.03	2.19		
PLACE1011017	5.6	3.78	22.57	25.64	35.47	20.97		
PLACE1011019	1.1	1.5	2.48	3.78	4.42	0.88		
PLACE1011026	4.17	4.93	6.23	6	5.19	2.74		
PLACE1011032	0.89	3.95	2.03	1.44	1.58	0.56		
PLACE1011041	1.07	4.13	1.03	1.69	1.22	1.43		
PLACE1011045	1.49	5.62	2.36	3.26	3.67	4.45		
PLACE1011046	0.83	5.25	1.79	2.57	3.17	1.87		
PLACE1011054	2.33	5.33	6.64	5.26	7.29	5.05		
PLACE1011056	5.78	5.43	16.22	14.56	19.78	15.67		
PLACE1011057	2	2.18	3.5	3.29	5.68	3.9		
PLACE1011059	0.93	1.37	1.56	1.74	2.96	1.79		

(1197)

特開2002-191363

2391

2392

PLACE1011066	4.49	5.74	6.76	5.38	7.72	5.49		
PLACE1011087	7.6	7.04	16.48	12.43	17.42	9.79		
PLACE1011090	2.98	6.14	6.74	3.36	4.13	3.26		
PLACE1011109	1.99	7.29	4.29	4.08	7.96	3.83		
PLACE1011114	1.62	4.4	3.13	3.33	4.68	3.29		
PLACE1011116	4.89	5.94	6.66	7.43	6.81	8.98		
PLACE1011122	0.93	2.52	1.84	2.1	1.61	1.64		
PLACE1011133	0.83	2.22	3.03	3.48	3.52	2.77		
PLACE1011134	12.47	15.29	66.86	44.95	68.68	51.65		
PLACE1011143	0.68	4.53	1.48	1.84	2.62	1.41		
PLACE1011146	0.91	5.93	1.74	1.97	3.23	2.36		
PLACE1011160	1.67	7.36	3.81	3.42	4.53	4.24		
PLACE1011165	1.77	2.34	3.39	2.15	3.39	3.8		
PLACE1011181	5.25	8.31	37.21	29.38	38.44	30.55		
PLACE1011185	2.47	4	3.57	4.66	5.15	4.8	*	+
PLACE1011186	13.16	12.8	16.45	21.18	25.69	28.83	*	+
PLACE1011203	1.08	4.64	2.19	1.75	2.94	1.96		
PLACE1011214	9.02	16.55	59.24	46.39	62.58	47.89		
PLACE1011219	1.41	4.91	3.6	3.26	4.96	2.9		
PLACE1011221	2.68	5.47	6.15	6.23	8.57	7.51		
PLACE1011229	1.38	4.2	2.69	2.43	2.43	3.09		
PLACE1011231	0.53	1.5	1.62	1.84	2.59	2.4		
PLACE1011236	5.69	7.9	43.53	33.48	56.77	44.25		
PLACE1011247	8.36	10.08	19.16	24.17	26.33	29.74	*	+
PLACE1011263	0.57	6.43	4.23	2.62	4.11	5.09		
PLACE1011273	0.72	3.62	1.17	2.01	1.64	1.72		
PLACE1011278	2.42	5	6.12	3.98	4.84	4.31		
PLACE1011289	2.73	5.84	7.57	6.34	6.13	6.08		
PLACE1011291	3.2	5.19	8.31	7.32	8.04	7.95		
PLACE1011296	0.93	2.45	1.94	1.76	2.63	2.05		
PLACE1011310	1.72	2.64	3.36	4.51	2.77	4.48		
PLACE1011311	1.8	3.97	6.33	7.8	9.34	6.73		
PLACE1011321	1.29	4.77	3.5	3.3	2.63	3.33		
PLACE1011325	0.63	4.2	1.84	2	2.74	1.59		
PLACE1011332	5.65	10.55	9.4	14.8	14.57	15.04	*	+
PLACE1011340	0.86	4.88	3.38	3.81	4.71	3.47		
PLACE1011353	5.39	5.53	8.39	10.06	8.58	4.43		
PLACE1011360	1.09	3.29	2.18	2.11	3.06	2.41		
PLACE1011364	4.88	5.69	7.92	6.34	4.57	4.57		
PLACE1011365	0.92	3.36	2.95	3.01	3.6	2.06		
PLACE1011371	9.37	10.69	63.06	56.51	87.51	64.33		
PLACE1011375	1.62	9.37	3.35	2.61	2.74	1.62		
PLACE1011386	3.91	12.96	4.18	6.88	5.51	5.93		
PLACE1011399	1.52	10.14	4.27	2.12	4.22	3.77		
PLACE1011406	1.25	2.05	1.81	5.03	3.26	2.38		
PLACE1011407	0.64	2.05	2.58	2.08	2.19	2.02		
PLACE1011419	2.83	3.2	6.02	6.23	4.18	3.72		
PLACE1011433	3.09	4.43	6.13	3.41	4.74	4.99		
PLACE1011440	0.77	5.81	2.48	2.1	2.24	2.27		
PLACE1011452	1.06	8.86	2.43	2.87	2.71	2.68		

(1198)

特開 2002-191363

2393

2394

PLACE1011465	0.09	8.53	1.96	1.04	1.68	0.85		
PLACE1011472	1.52	7.67	3.98	0.97	2.87	1.59		
PLACE1011477	11.18	12.29	54.35	31.86	55.86	45.67		
PLACE1011478	1.24	1.64	3.11	5.03	3.02	3.01		
PLACE1011492	2.24	3.54	5.01	5.45	6.74	5.27		
PLACE1011498	0.57	2.77	0.98	1.18	1.69	1.15		
PLACE1011501	0.49	4.67	1.8	4.87	5.63	7.5		
PLACE1011503	0.44	5.37	0.79	0.5	0.82	0.35		
PLACE1011509	1.1	7.8	3.38	4.62	5.08	5.74		
PLACE1011514	5.86	11.61	11.98	13	17.7	13.17		
PLACE1011516	10.37	13.29	18.32	18.32	8.08	3.63		
PLACE1011520	0.34	0.95	1.34	1.3	1.73	1.01		
PLACE1011538	52.87	99.27	185.04	129.45	86.87	63.9		
PLACE1011555	0.87	2.88	2.83	1.55	2.03	1.48		
PLACE1011561	3.2	4.53	1.98	6.81	5.31	3.08		
PLACE1011563	1.35	4.74	3.49	2.52	2.64	2.87		
PLACE1011567	1.04	4.94	3.36	2.75	4.19	2.77		
PLACE1011569	0.32	4.35	2.77	2.38	2.46	2.28		
PLACE1011576	3.25	1.88	7.94	7.85	9.1	8.03		
PLACE1011586	3.24	2.5	6.22	4.43	4.35	2.94		
PLACE1011635	1.85	2.56	4.53	9.96	12.43	8.71	**	+
PLACE1011641	0.43	2.9	0.9	1.71	1.18	1.19		
PLACE1011642	5.05	6.96	10.37	12.13	10.86	10.09		
PLACE1011643	1.29	3.69	1.14	2.38	2.28	2.14		
PLACE1011646	8.68	12.8	30.53	39.05	54.16	37.76	*	+
PLACE1011649	1.35	5.7	3.88	4.5	4.61	5.46		
PLACE1011650	1.15	1.45	2.54	2.47	3.24	3.75		
PLACE1011661	1.02	2.26	2.8	3.95	5.92	3.46		
PLACE1011664	2.21	3.18	3.99	5.31	3.93	1.73		
PLACE1011672	0.88	4.14	0.72	2.69	3.57	2.04		
PLACE1011675	0.51	2.31	1.32	1.66	0.99	0.83		
PLACE1011682	2.04	4.56	2.23	2.22	4.03	2.12		
PLACE1011708	1.1	5.89	3.8	5.5	8.12	4.66		
PLACE1011719	1.07	4.58	1.66	3.55	3.88	3.03		
PLACE1011725	1.23	1.19	2.72	3.73	5.19	4.3	*	+
PLACE1011729	0.86	1.03	1.8	2.38	3.26	1.22		
PLACE1011741	2.36	3.67	3.64	4.16	2.52	4.23		
PLACE1011749	1.58	3.89	4.09	3.49	4.85	3.27		
PLACE1011757	20.92	30.53	55.88	56.6	55.88	49.59		
PLACE1011762	0.4	4.34	2.69	3.91	2.14	2.3		
PLACE1011778	0.51	4.39	1.99	1.34	2.02	1.34		
PLACE1011783	2.59	4.63	5.46	4.8	8.41	5.55		
PLACE1011795	0.74	1.28	2.24	1.47	1.8	1.22		
PLACE1011810	9.28	10.82	19.51	13.32	17.73	15.47		
PLACE1011824	5.38	8.17	38.05	25.52	42.89	16.75		
PLACE1011825	10.61	16.39	22.6	17.63	18.92	16.07		
PLACE1011835	24.64	32.67	47.67	32.09	26.75	37.23		
PLACE1011836	18.11	18.97	31.43	33.14	47.23	13.95		
PLACE1011847	2.67	6.74	6.42	5.34	7.18	5.84		
PLACE1011855	0.9	6.06	3.53	3.23	4.47	3.49		

(1199)

特開 2002-191363

2395

2396

PLACE1011858	5.83	7.44	9.37	6.88	6.25	8.35		
PLACE1011874	1.35	3.14	3.55	4.18	5.28	4.65		
PLACE1011875	0.57	2.29	1.11	2.66	2.48	2.64		
PLACE1011877	3.8	5.03	4.4	7.67	6.8	9.97	*	+
PLACE1011891	0.17	3.81	1.31	1.13	1.34	1.26		
PLACE1011896	-0.1	5.22	1.45	-0.19	1.07	0.32		
PLACE1011920	0.21	5.87	1.04	1.1	1.92	1.76		
PLACE1011922	2.4	4.18	4.72	3.72	2.88	3.98		
PLACE1011923	3.42	4.82	7.51	8.09	5.38	12.28		
PLACE1011937	3.16	2.24	3.76	3.81	5.58	4.64		
PLACE1011939	14.93	17.81	26.01	41.75	45.05	47.88	**	+
PLACE1011940	6.13	7.58	12.68	5.73	7	7.23		
PLACE1011962	3.28	7.83	7.35	6.39	8.38	8.01		
PLACE1011964	0.39	5.05	1.66	1.04	1.95	1.46		
PLACE1011978	1.55	4.65	3.35	4.48	5.91	5.14		
PLACE1011980	2.1	4.62	6.07	3.95	4.91	5.35		
PLACE1011981	5.77	7.28	38.2	27.22	35.24	34.99		
PLACE1011982	0.83	3.1	2.23	1.82	2.59	1.57		
PLACE1011995	0.81	3.4	3.73	2.51	3.59	2.86		
PLACE1012023	1.38	5.37	1.87	2.09	2.46	1.91		
PLACE1012026	1.95	5.72	4.23	6.08	9.51	9.8		
PLACE1012031	2.49	5.81	4.54	4.34	6.35	4.03		
PLACE2000003	1.18	3.64	6.86	7.38	8.12	8.92		
PLACE2000005	1.16	2.41	2.16	2.76	2.03	1.89		
PLACE2000006	2.52	4.13	15.6	11.34	16.58	13.4		
PLACE2000007	0.96	4.85	4.24	3.94	5.13	3.33		
PLACE2000011	1.72	3.27	3.34	4.3	5.06	3.62		
PLACE2000014	4.04	5.93	23.94	27.19	30.87	30.97	*	+
PLACE2000015	1.27	4.79	3.52	2.77	3.31	2.69		
PLACE2000017	0.48	4.78	2.15	2.65	2.56	2.45		
PLACE2000021	1.99	5.06	4.09	3.72	6.24	5.61		
PLACE2000022	1.8	2.8	4.31	3.35	4.64	2.71		
PLACE2000030	9.37	11.08	71.38	55.43	60.32	35.97		
PLACE2000032	1.23	2.89	3.51	3.53	3.24	2.32		
PLACE2000033	3.29	6.3	12.11	12.49	14.25	9.37		
PLACE2000034	0.6	4.3	1.92	1.79	2.19	2.44		
PLACE2000039	2.75	8.06	3.92	6.27	8.01	4.81		
PLACE2000043	7	11.08	22.94	19.27	26.58	17.38		
PLACE2000044	0.63	6.92	1.12	1.8	1.71	1.29		
PLACE2000047	0.84	5.25	5.15	5.4	6.46	7.32		
PLACE2000050	1.48	2.68	4.78	2.92	2.98	1.2		
PLACE2000061	0.47	2.07	1.17	0.95	1.34	0.25		
PLACE2000062	1.99	2.83	4.12	4.89	4.4	3.09		
PLACE2000072	0.78	2.45	1.57	1.85	1.62	1.69		
PLACE2000073	0.89	5.86	2.86	2.8	2.61	3.07		
PLACE2000097	8.54	19.93	23.93	27.69	35.36	26.36		
PLACE2000100	1.87	7.79	3.53	4.23	6.29	4.22		
PLACE2000103	1.03	7.44	2.25	3.51	5.17	4.38		
PLACE2000106	1.53	2.42	4.19	5.71	3.29	4.01		
PLACE2000111	2.05	3.17	4.37	4.07	6.6	4.75		

(1200)

特開 2002-191363

2397

2398

PLACE2000115	0.3	2.06	0.75	0.31	1.29	0.97		
PLACE2000118	10.15	17.04	21.09	13.73	15.21	22.01		
PLACE2000124	10.14	17.83	62.13	53.2	98.37	62.96		
PLACE2000132	0.06	6.26	1.48	0.99	1.5	1.56		
PLACE2000136	0.55	7.94	0.91	1.47	1.37	1.02		
PLACE2000137	0.96	4.46	2.4	2.65	4.12	3.14		
PLACE2000140	2.91	5.24	14.34	13.2	12.08	9.43		
PLACE2000147	1.49	1.52	2.83	1.06	0.97	1.12		
PLACE2000153	0.3	3.44	2.15	1.69	2.45	1.95		
PLACE2000164	0.66	2.78	1.13	2.66	1.62	1.62		
PLACE2000170	1.54	6.18	4.69	5.26	9.09	6.24		
PLACE2000172	0.33	4.34	2.15	1.25	1.93	2.43		
PLACE2000173	0.92	4.97	3.37	3.33	3.71	2.74		
PLACE2000174	1.17	4.68	2.43	1.85	2.89	2.05		
PLACE2000176	1.22	1.57	2.58	3.28	2.27	1.42		
PLACE2000187	1.01	2.08	2.55	3.45	3.66	2.19		
PLACE2000216	7.03	9.28	11.47	14.09	9.13	3.68		
PLACE2000219	0.69	4.02	2.72	3.55	3.58	2.3		
PLACE2000221	2.49	6.81	6.53	7.22	9.33	8.56		
PLACE2000223	0.72	3.2	1.71	1.16	1.05	0.48		
PLACE2000231	1.02	3.97	3.11	3.43	2.61	2.47		
PLACE2000235	1.82	5.27	6.17	6.02	7.45	6.97		
PLACE2000246	1.93	2	6.06	4.58	5.09	3.93		
PLACE2000264	0.67	1.39	1.85	2.45	3.74	3	*	+
PLACE2000274	0.65	2.4	2.12	3.09	4.11	2.1		
PLACE2000287	0.81	4.44	1.49	2	2.59	1.34		
PLACE2000296	1.01	4.56	1.55	2.5	3.16	1.69		
PLACE2000302	1.34	4.67	2.86	3.52	3.35	3.45		
PLACE2000305	3.09	6.65	5.42	6.3	7.15	5.02		
PLACE2000317	0.92	6.34	2.26	3.15	2.95	2.4		
PLACE2000324	1.19	1.25	3.09	4.2	5.84	3.3		
PLACE2000334	3.52	5	6.6	7.33	8.12	5.88		
PLACE2000335	1.47	4.35	4.24	5.68	6.25	4.76		
PLACE2000340	0.64	3.47	1.63	1.58	1.52	1.65		
PLACE2000341	4.21	7.87	28.81	18.94	32.45	19.16		
PLACE2000342	2.07	5.11	4.32	4.84	5.82	4.49		
PLACE2000347	1.24	5.26	5.58	7.83	6.22	7.55		
PLACE2000357	8.49	13.56	15.35	17.83	18.98	21.24	*	+
PLACE2000358	2.87	3.65	8.67	4.88	7.63	4.37		
PLACE2000359	1.27	1.79	4.45	3.28	6.65	3.61		
PLACE2000366	1.93	3.14	3.22	3.99	5.6	4.17		
PLACE2000371	4.29	5.2	6.08	5.95	9.06	7.32		
PLACE2000373	1.91	4.8	5.98	5.69	6.29	4.19		
PLACE2000374	1.86	5.17	2.78	1.62	2.79	1.49		
PLACE2000379	0.34	4.85	1.32	1.28	0.92	0.04		
PLACE2000386	39.29	43.92	84.66	87.53	104.55	76.56		
PLACE2000388	1.96	3.35	3.89	2.78	3.48	3.49		
PLACE2000392	33.29	39.2	59.56	42.5	52.12	58.24		
PLACE2000394	1.26	3.27	3.01	5.69	4.35	4.34	*	+
PLACE2000398	0.73	3.88	2.36	2.03	3.59	1.35		

(1201)

特開2002-191363

2399

2400

PLACE2000399	3.7	6.82	7.01	7.15	7	6.79		
PLACE2000402	2.15	6.88	3.84	2.86	3.68	3.9		
PLACE2000404	5.2	9.96	10.67	10.03	11.04	6.47		
PLACE2000411	3.21	7.2	5.21	5.27	5.73	6.68		
PLACE2000418	0.73	2.28	2.41	2.22	3.07	2.37		
PLACE2000419	0.99	2.32	2.54	4.95	4.55	3.29	*	+
PLACE2000425	1.26	3.98	3.11	4.28	4.81	5.2		
PLACE2000427	0.7	5.13	3.27	2.54	3.04	2.47		
PLACE2000433	0.77	7.05	2.6	2.33	3.09	2.46		
PLACE2000435	0.48	5.19	1.49	1.69	1.63	1.5		
PLACE2000438	1.61	4.74	3.66	2.33	2.81	3.15		
PLACE2000450	3.01	4.38	5.67	6.51	7.39	5.63		
PLACE2000455	0.24	2.62	1.24	1.65	2	1.82		
PLACE2000458	0.38	3.3	1.81	1.06	2.7	1.24		
PLACE2000464	2.15	4.91	5.3	7.43	9.68	8.83	*	+
PLACE2000465	1.43	6.72	6	6.51	8.27	6.31		
PLACE2000473	120.94	179.35	328.3	214.7	297.75	279.74		
PLACE2000477	0.43	3.87	1.34	1.13	2.22	0.97		
PLACE3000004	2.22	4.63	6.39	5.27	7.51	5.2		
PLACE3000009	19.91	19.71	105.63	77.3	140.99	92.95		
PLACE3000020	10.03	9.03	49.6	36.74	46.52	23.82		
PLACE3000029	6.59	9.63	24.88	14.88	18.47	20.04		
PLACE3000038	0.52	2.37	2.47	1.44	2.4	2.05		
PLACE3000052	5.13	7.95	23.92	25.01	29.61	24.94		
PLACE3000059	0.57	5	2.42	0.75	2.8	1.27		
PLACE3000067	2.51	5.79	7.44	5.66	8.53	7.75		
PLACE3000069	1.95	5.61	3.58	5.24	3.79	4.55		
PLACE3000070	2.57	5.57	9.04	9.5	10.42	10.57		
PLACE3000103	3.85	7.84	11.87	6.6	8.32	4.37		
PLACE3000119	1.59	2.74	3.15	3.24	3.67	2.95		
PLACE3000121	7.58	8.44	38.1	30.63	42.28	32.64		
PLACE3000124	1.53	4.54	5.95	6.35	7.75	7.18		
PLACE3000135	0.69	5.46	1.1	0.76	0.9	0.59		
PLACE3000136	0.77	10.46	4.46	2.12	2.77	2.01		
PLACE3000142	0.7	9.94	1.75	1.53	2.76	1.13		
PLACE3000145	8.69	17.55	55.33	42.85	49.12	39.96		
PLACE3000147	15.7	12.92	39.97	64.96	54.76	30.34		
PLACE3000148	0.7	2.08	1.48	0.82	1.4	1.38		
PLACE3000154	0.48	1.86	0.67	1.5	0.87	0.44		
PLACE3000155	1.28	4.26	2.53	3.97	4.76	3.04		
PLACE3000156	1.11	7.96	2.07	1.96	3.4	2.81		
PLACE3000157	0.92	8.37	1.75	2.06	3.02	2.72		
PLACE3000158	1.73	8.82	5.05	3.81	5.63	4.77		
PLACE3000160	8.11	15.22	19.5	33.66	34.71	33.78	**	+
PLACE3000169	2.15	2.65	6.87	4.3	5.11	5.12		
PLACE3000181	1.06	2.14	3.94	3.22	2.62	3		
PLACE3000194	0.31	2.83	1.77	1.89	2.58	2.42		
PLACE3000197	1.18	2.66	2.09	2.42	2.46	2.21		
PLACE3000199	0.22	4.8	1.63	0.55	0.86	0.43		
PLACE3000205	11.79	17.95	57.49	51.4	63.83	42.63		

(1202)

特開2002-191363

2401

2402

PLACE3000207	3.37	6.91	7.13	6.7	8.92	6.23		
PLACE3000208	2.26	4.96	2.66	5.28	5.97	5.84		
PLACE3000213	4.79	5.55	10.8	5.12	5.16	5.05		
PLACE3000215	1.88	5.02	5.71	4.74	5	3.03		
PLACE3000218	0	1.63	1.18	0.97	0.62	0.31		
PLACE3000220	1.96	3.55	4.58	6.74	7.52	6.69	*	+
PLACE3000221	14.42	25.34	40.15	43.8	51.16	36.99		
PLACE3000225	1.15	4.68	3.11	2.11	2.67	1.28		
PLACE3000226	1.37	5.65	5.16	3.78	7.42	3.68		
PLACE3000230	0.83	3.46	1.36	2	2.8	1.73		
PLACE3000231	1.31	1.97	2.37	4.86	3.95	4.12	**	+
PLACE3000235	1.12	1.75	3.89	3.95	4.21	3.39		
PLACE3000242	2.6	5.11	9.24	9.46	10.97	8.29		
PLACE3000244	1.05	3.2	1.81	1.85	1.81	0.64		
PLACE3000253	0.7	3.75	1.64	2.67	2.11	1.27		
PLACE3000254	2.5	4.75	4.04	6.19	6.09	5.75	*	+
PLACE3000271	2.67	6.06	6.81	10.96	10.99	9.5	*	+
PLACE3000276	1.1	5.78	2.27	1.48	1.9	1.78		
PLACE3000304	5.55	4.69	10.81	11.19	11.49	10.5		
PLACE3000309	0.43	1.67	1.87	2.43	2.94	2.78	*	+
PLACE3000310	2.19	2.19	3.73	4.84	4.81	3.4		
PLACE3000320	1.02	3.65	1.8	2.54	2.37	2.32		
PLACE3000322	1.31	4.23	6.63	7.5	7.8	6.09		
PLACE3000330	24.05	24.44	41.08	31.87	35.83	29.17		
PLACE3000331	1.21	5.86	4.14	4.34	5.7	4.31		
PLACE3000336	2.61	6.99	4.42	4.24	5.72	5.11		
PLACE3000339	7.36	5.1	11.41	16.25	18.28	17.37	**	+
PLACE3000341	1.65	1.32	2.41	4.08	4.35	3.65	**	+
PLACE3000350	5.88	6.4	12.86	15.45	18.5	15.41	*	+
PLACE3000352	1.54	3.88	2.13	2.37	2.25	1.71		
PLACE3000353	5.38	9.72	11.8	19.12	22.98	15.5	*	+
PLACE3000362	0.62	4.92	4.72	3.61	5.33	3.39		
PLACE3000363	2.19	5.13	2.32	1.89	3.28	2.07		
PLACE3000365	1.34	6.11	3.37	3.34	4.05	2.12		
PLACE3000373	0.89	1.52	3.66	2.93	6.08	2.3		
PLACE3000374	1.07	1.85	2.91	2.72	2.99	2.15		
PLACE3000387	0.31	3.32	1.04	1.24	1.65	1.29		
PLACE3000388	1.18	3.22	1.94	2.76	3.49	2.22		
PLACE3000399	2.12	4.66	6.28	7.42	9.84	6.05		
PLACE3000400	3.08	5.44	11.87	7.97	10.77	7.82		
PLACE3000401	7.52	11.42	18.59	22.61	29.55	23.4	*	+
PLACE3000402	1.79	3.21	3.4	2.19	1.74	1.79		
PLACE3000405	3.37	3.74	5.82	5.54	7.22	6.01		
PLACE3000406	2.1	2.91	3.11	3.48	3.68	2.42		
PLACE3000413	1.18	2.72	2.69	1.71	2.06	1.52		
PLACE3000416	1.05	4.03	4.04	3.43	2.67	2.72		
PLACE3000425	1.21	6.27	4.33	3.98	6.36	3.92		
PLACE3000437	4.79	10.85	29.89	16.69	25.26	19.14		
PLACE3000455	2.97	8.07	10.62	8.97	10.39	7.91		
PLACE3000475	16.52	19.2	47.35	40.22	39.77	34.21		

(1203)

特開 2002-191363

2403

2404

PLACE3000477	5.44	4.79	5.56	8.05	5.52	8.42		
PLACE4000003	0.38	2.97	1.61	3.14	2.33	2.31		
PLACE4000008	15.19	11.38	16.76	13.05	14.26	8.84		
PLACE4000009	1.17	6.19	3.39	3.93	3.37	1.82		
PLACE4000014	1.31	5.12	1.77	2.16	3.03	2.19		
PLACE4000029	6.33	8.48	35.37	23.93	32.21	24.25		
PLACE4000034	2.27	6.24	5.22	6.46	6.52	4.91		
PLACE4000049	3.39	3.35	5.21	3.85	5.82	4.86		
PLACE4000052	1.41	3.36	2.2	2.62	2.64	2.02		
PLACE4000062	1.6	4.94	5.06	4.25	5.06	3.71		
PLACE4000063	2.59	6.87	5.19	4.86	4.81	3.73		
PLACE4000089	1.52	6.31	3.35	2.81	3.91	2.84		
PLACE4000093	0.44	5.6	1.61	1.28	1.65	1.98		
PLACE4000100	2.72	6.13	4.75	4.33	3.62	3.94		
PLACE4000103	0.63	4.48	5.64	4.4	5.67	2.9		
PLACE4000106	3.2	5.33	6.63	7.1	5.13	7.21		
PLACE4000128	1.93	3.97	4.88	4.15	4.96	4.4		
PLACE4000129	0.74	3.26	1.64	1.57	2.11	1.78		
PLACE4000131	7.14	10.85	41.43	32.45	41.08	31.22		
PLACE4000147	0.34	3.65	0.54	0.45	0.93	0.61		
PLACE4000156	2.47	6.08	8.06	7.83	13.47	9.07		
PLACE4000175	0.72	4.08	1.48	0.98	0.91	0.84		
PLACE4000190	14.55	18.47	70.34	49.15	74.82	60.76		
PLACE4000192	1.3	2.27	3.6	2.36	2	1.25		
PLACE4000206	5.35	6.65	12.44	7.13	7.1	6.02		
PLACE4000211	3.34	4.64	22.23	11.68	12.35	13.44		
PLACE4000214	0.86	3.61	2.68	2.08	2.53	1.69		
PLACE4000222	0.93	5.28	4.36	4.13	4.75	3.5		
PLACE4000223	0.46	4.51	1.79	1.37	1.22	0.38		
PLACE4000229	1.9	5.79	2.11	2.81	3.36	3.48		
PLACE4000230	1.11	5.89	6.51	3.61	6.81	5.15		
PLACE4000233	1.26	3.02	5.66	2.92	2.98	3.51		
PLACE4000239	2.35	3.68	4.17	4.19	3.97	3.35		
PLACE4000247	0.52	2.37	3.38	2.64	3.1	2.35		
PLACE4000250	1.18	3.24	2.35	3.33	3.68	2.8		
PLACE4000252	1.06	4.99	2.25	1.92	1.75	1.64		
PLACE4000259	4.42	11.95	18.1	14.47	22.09	14.02		
PLACE4000261	0.87	10.29	1.07	2.03	1.9	1.12		
PLACE4000264	15.86	24.96	36.9	11.96	21.82	22.51		
PLACE4000269	3.48	3.71	7.95	4.62	4.55	2.85		
PLACE4000270	0.43	1.42	1.87	1.75	1.83	0.59		
PLACE4000281	17.84	20.97	44.05	32.93	28.37	28.87		
PLACE4000300	0.67	2.06	2.04	3.21	2.88	3.58	*	+
PLACE4000320	1.33	5.86	3.1	2.84	5.32	3.21		
PLACE4000323	1.63	7.43	5.13	4.03	4.65	4.82		
PLACE4000326	1.8	10.98	5.67	5.72	8.73	5.59		
PLACE4000344	0.22	5.75	2.62	1.66	1.6	1.18		
PLACE4000347	4.7	3.82	13.93	16.83	16.75	17.36	*	+
PLACE4000354	3.18	6.29	10.68	5.17	2.81	2.79		
PLACE4000367	0.79	2.97	1.71	0.87	1.3	1.38		

(1204)

特開 2002-191363

2405

2406

PLACE4000369	1.35	3.97	2.36	1.99	1.96	0.82
PLACE4000379	2.44	6.66	5.44	5.94	7.55	5.07
PLACE4000387	0.88	5.86	2.11	1.28	0.84	1.12
PLACE4000392	0.42	5.58	1.32	1.81	1.02	1.63
PLACE4000399	10.99	17.08	75.17	59.11	80.22	58
PLACE4000401	0.72	0.7	1.53	1.17	0.83	1.4
PLACE4000403	3.15	4.13	8.51	5.29	6.38	5.87
PLACE4000411	2.22	2.28	4	2.27	2.6	1.82
PLACE4000415	0.7	3.55	2.8	1.16	1.86	0.78
PLACE4000416	25.49	29.13	33.54	23.65	21.92	24.83
PLACE4000424	1.61	5.59	3.33	3.27	3.92	2.51
PLACE4000431	3.89	7.39	21.01	17.68	28.21	16.79
PLACE4000443	0.07	4.33	2.15	1.52	2.83	1.14
PLACE4000445	3.94	5.43	9.98	7.62	6.99	6.27
PLACE4000450	2.99	3.65	23.28	15.51	24.53	16.04
PLACE4000455	5.18	7.39	9.55	8	7.21	4.63
PLACE4000465	1.39	4.34	3.26	4.15	6.07	4.34
PLACE4000466	120.96	98.04	201.25	113.83	170.96	145.31
PLACE4000472	3.12	9.6	10.17	10.92	13.21	9.22
PLACE4000487	3.18	7.83	16.5	14.66	16.62	15.05
PLACE4000489	0.93	4.69	3.41	1.95	3.88	1.69
PLACE4000494	1.15	1.6	4.07	2.74	3.1	2.08
PLACE4000502	6.3	5.39	10.92	11.65	15.08	6.37
PLACE4000521	2.5	3.44	16.06	12.78	20.63	11.2
PLACE4000522	5.07	6.17	9.07	12.43	8.68	14.11
PLACE4000537	0.98	4.28	1.27	1.67	1.61	1.22
PLACE4000548	1.99	5.69	2.46	3.04	3.68	2.32
PLACE4000558	0.87	6.72	1.97	3.15	2.41	2.15
PLACE4000581	2.1	7.22	7.04	3.9	5.96	5.44
PLACE4000590	0.4	0.61	0.15	0.4	0.81	-0.25
PLACE4000593	2.94	2.98	5.22	4.44	5.82	3.83
PLACE4000612	0.68	3.33	3.33	1.5	3.02	2.74
PLACE4000638	1.25	4.24	0.84	1.2	1.44	1.58
PLACE4000650	0.82	4.67	1.02	1.43	1.11	1.16
PLACE4000651	2.42	6.4	7.48	5	7.01	6.07
PLACE4000654	0.98	5.7	2.47	1.35	2.48	1.47
PLACE4000670	0.5	4.06	2.92	0.76	1.29	0.67
PLACE4000685	6.35	8.68	13.83	13.46	14.26	13.77
PLACE4000687	0.37	3.02	1.11	2.2	1.4	1.12
PLACE5000003	1.1	2.74	3.31	3.21	3.55	3.07
PLACE5000005	12.43	16.53	27.36	24.54	24.57	24.76
PLACE5000019	0.4	4.15	1.13	0.59	1.89	0.79
PLACE5000021	0.74	4.59	1.61	0.39	0.93	0.32
PLACE5000022	1.2	6.11	2.25	3.17	2.76	2.09
PLACE5000024	1.77	2.58	2.27	2.92	3.39	3.84
PLACE5000036	1.81	3.24	3.11	2.41	3.19	2.84
PLACE5000059	14.41	17.79	26.55	25.98	30.03	34.87
PLACE5000076	1.41	3.61	2.22	4.04	3.96	2.54
PLACE5000117	7.44	12.48	15.66	16.87	18.78	20.64
PLACE5000143	0.85	6.45	2.11	1.67	2.85	2.73

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(1205)

特開2002-191363

2407

2408

PLACE5000152	0.42	4.49	1.23	1.61	1.95	1.57		
PLACE5000154	18.23	23.5	45.06	21.81	25.65	31.8		
PLACE5000155	3.35	2.81	5.51	3.94	2.78	4.87		
PLACE5000165	3.78	4.4	6.67	4.51	5.99	5.82		
SKNMC1000004	9.7	11.62	16.77	10.19	12.16	13.96		
SKNMC1000011	1.82	8.58	4.12	5.89	3.95	6.77		
SKNMC1000013	0.51	6.69	1.13	1.21	2.14	1.44		
SKNMC1000014	1.28	4.18	3.22	3.77	6.37	3.96		
SKNMC1000018	3.42	5.19	5.25	5.51	5.68	3.44		
SKNMC1000020	0.95	4.03	3.46	3.6	4.68	4.56		
SKNMC1000046	2	3.17	3.48	3.95	3.26	2.55		
SKNMC1000050	4.99	8.04	10.32	5.4	6.28	6.12		
SKNMC1000062	9.79	12.6	20.18	19.2	15.42	18.73		
SKNMC1000075	1.45	4.3	2.01	1.98	1.89	2.92		
SKNMC1000082	1.12	4.39	2.13	1.85	1.78	2.39		
SKNMC1000091	4.54	7.52	7.95	11.74	12.86	12.77	**	+
SKNMC1000099	0.33	4.29	1.98	1.32	0.65	1.18		
SKNMC1000104	1.13	4.24	3.45	1.47	3.14	2.43		
SKNMC1000113	0.97	1.83	1.2	1.74	2.63	0.89		
SKNMC1000119	1.73	2.64	5.07	4.48	5.34	4.67		
SKNMC1000142	0.04	2.87	0.99	1.27	0.75	1		
SKNMC1000170	0.91	4.75	2.34	1.71	1.49	1.11		
SKNMC1000178	3.02	8.39	7.08	5.77	9.65	9.02		
SKNMC1000194	0.63	9.82	1.51	0.61	1.73	1.3		
SKNMC1000198	1.35	11.01	3.33	2.65	2.1	2.88		
SKNMC1000225	1.35	6.44	2.97	2.39	3.4	3.26		
SKNMC1000249	0.49	2.14	0.75	0.57	0.51	0.52		
SPLN1000007	0.74	2.15	2.11	1.7	2.26	1.99		
SPLN1000012	0.39	1.9	1.72	1.19	0.8	0.84		
SPLN1000014	1.78	4.4	4.9	5.75	4.33	3.99		
SPLN1000036	4.95	11.64	24.32	20.56	27.73	21.68		
SPLN1000059	0.04	6.69	1.06	0.91	1.79	1.47		
SPLN1000068	1.68	10.81	5.71	5.79	5.17	5.64		
SPLN1000072	1	8.5	4.7	2.82	3	2.21		
SPLN1000101	20.01	18.4	45.64	29.93	25.24	12.63		
SPLN1000108	0.56	1.54	0.98	0.75	1.11	0.76		
SPLN1000113	1.33	2.27	3.04	2.72	4.13	3.04		
SPLN1000114	2.97	4.19	6.03	3.59	4.76	6.32		
SPLN1000132	0.85	4	1.72	2.25	2.67	1.99		
SPLN1000135	3.13	8.76	14.93	11.12	15.28	10.52		
SPLN1000136	12.41	21.47	15.14	20.24	27.48	21.8		
SPLN1000141	2.26	7.07	10.79	4.03	5.41	4.51		
SPLN1000164	2.49	3.79	8.58	3.98	5.88	7.61		
SPLN1000166	0.4	2.9	2.96	1.67	1.19	1.68		
SPLN1000175	2.16	4.48	6.1	5.65	4.12	4.15		
SPLN1000182	0.98	2.66	0.23	0.83	0.6	0.67		
SPLN1000185	3.41	8.49	8.54	11.38	10.43	11.95		
THYMU1000004	10.22	14.07	20.43	22.34	22.76	23.6		
THYMU1000009	9.48	10.13	14.9	13.48	23.86	22.1		
THYMU1000015	8.87	10.42	16.18	19.25	22.21	20.8	*	+

(1206)

特開2002-191363

2409

2410

THYMU1000016	6.24	5.96	13.03	10.3	8.45	9.38
THYMU1000023	0.77	1.86	3.6	5.22	3.68	3.6
THYMU1000034	0.16	1.77	1.8	0.79	0.88	0.14
THYMU1000035	0.62	2.8	0.97	1.17	0.95	1.31
THYMU1000037	1.53	4.15	2.11	2.06	2.81	1.45
THYMU1000042	5.97	10.24	12.23	12.03	13.98	13.28
THYMU1000047	2.72	6.03	6.72	6.04	7.77	7.23
THYMU1000080	0.56	4.31	2.6	3.26	1.85	2.11
THYMU1000094	2.77	3.47	7.91	9.17	8.35	4.55
THYMU1000109	17.28	14.34	111.37	98.05	142.29	93.04
THYMU1000127	2.75	5.95	10.76	8.18	9.98	6.74
THYMU1000130	2.5	4.4	4.55	6.69	6.07	4.94
THYMU1000137	3.53	7.18	10.26	12.67	18.55	13.05
THYMU1000146	4.37	8.38	6.52	8.29	7.46	7.74
THYMU1000159	5.43	9.51	16.37	12.4	15.15	13.27
THYMU1000163	5.85	12.26	37.58	45.53	58.37	36.93
THYMU1000167	2.39	3.02	4.73	4.89	6.79	3.97
THYMU1000186	0.69	1.05	1.45	1.31	2.45	0.66
THYRO1000017	0.94	3.45	2.54	2.02	3.54	2.11
THYRO1000026	1.56	5.63	4.02	3.96	4.82	3.36
THYRO1000034	0.49	4.16	1.59	1.99	2	1.82
THYRO1000035	0.86	4.84	1.34	2.29	2.48	2.11
THYRO1000036	0.93	8.32	4	3.08	4.36	5.59
THYRO1000040	2.58	7.02	4.76	4.66	4.83	4.93
THYRO1000061	2.01	1.91	3.07	3.53	3.8	2.61
THYRO1000067	1.98	2.8	5.12	3.37	4.14	3.3
THYRO1000070	1.26	2.09	3.59	2.65	3.85	3.45
THYRO1000072	1.33	3.37	4.22	2.54	4.08	2.06
THYRO1000084	8.07	12.69	22.39	2.99	5.61	4.42
THYRO1000085	1.44	5.66	3.99	2.42	3.86	2.85
THYRO1000086	-0.05	5.46	1.74	0.89	1.18	1.15
THYRO1000087	0.72	3.86	1.01	0	0.58	0.17
THYRO1000092	2.32	5.1	4.66	3.75	4.43	4.5
THYRO1000093	0.35	3.24	0.83	1.54	1.27	0.95
THYRO1000099	0.45	2.53	2.73	2.8	1.67	2.39
THYRO1000107	0.5	2.95	2.7	2.86	3.22	2
THYRO1000111	0.85	4.58	1.78	1.4	2.06	2.36
THYRO1000121	1.33	5.72	2.52	1.94	2.4	2.95
THYRO1000124	0.27	5.55	0.64	0.86	0.89	0.64
THYRO1000129	0.36	2.1	0.11	0.94	1.11	0.92
THYRO1000130	1.82	3.11	3.13	3.85	3.01	2.39
THYRO1000132	2.4	3.62	9.43	11.14	6.99	6.26
THYRO1000134	1.5	4.07	3.22	4.06	3.65	3.73
THYRO1000144	1.72	4.78	3.15	7.87	7.09	2.33
THYRO1000155	1.6	4.1	1.45	1.77	1.9	2.23
THYRO1000156	1.13	6.53	3.62	2.45	4.29	2.58
THYRO1000163	3.62	8.42	5.28	4.76	6.63	2.24
THYRO1000173	1.19	4.45	2.26	3.33	1.36	2.75
THYRO1000186	1.98	3.24	7.86	6.91	6.84	6.35
THYRO1000187	2.7	3.58	5.3	4.92	6.24	5.22

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(1207)

特開2002-191363

2411

2412

THYRO1000190	1.12	3.32	2.94	3.73	4.55	2.71		
THYRO1000196	0.3	5.28	0.81	0.66	1.21	0.52		
THYRO1000197	2.05	7.28	4.69	4.08	6.24	3.89		
THYRO1000199	0.76	6.28	4.13	1.93	2.08	1.98		
THYRO1000206	8.47	6.92	9.25	8.44	11.6	7.5		
THYRO1000221	1.9	3.17	4.42	4.02	5.87	4.54		
THYRO1000222	3.65	4.26	4.23	4.68	4.96	4.93	*	+
THYRO1000228	0.81	3.67	2.85	2.24	3.04	2.94		
THYRO1000241	1.76	3.7	6.29	4.62	5.54	4.01		
THYRO1000242	0.63	4.16	4.46	2.49	2.56	2.62		
THYRO1000246	1.61	5.5	3.9	3.43	4.7	3.91		
THYRO1000253	1.07	4.05	1.73	1.99	3.35	2.31		
THYRO1000270	1.15	5.12	1.39	1.22	2.5	1.26		
THYRO1000279	0.42	2.84	0.25	0.65	1.01	0.58		
THYRO1000285	2.75	4.65	7.31	7.03	7.75	4.88		
THYRO1000288	7.76	7.59	11.77	5.68	5.07	7.22		
THYRO1000296	4.18	6.04	6.22	7.4	11.24	8.96	*	+
THYRO1000320	1.54	5.83	4.97	3.65	4.45	3.34		
THYRO1000322	1.1	5.48	2.48	1.76	3.93	1.76		
THYRO1000327	1.75	7.69	4.77	6.21	5.23	4.41		
THYRO1000343	2.5	6.12	5.35	5.06	5.04	6.13		
THYRO1000345	1.36	7.34	11.92	7.82	5.84	9.49		
THYRO1000358	1.82	3.39	3.08	1.92	2.32	1.54		
THYRO1000368	0.76	2.39	2.73	1.43	2.82	0.58		
THYRO1000375	3.2	7.03	4.79	7.38	6.09	9.77		
THYRO1000381	0.92	2.88	2.19	3.87	3.11	2.74		
THYRO1000387	0.98	6.66	3.22	2.53	3.56	2.51		
THYRO1000394	1.31	9.88	4.59	4.29	5.19	2.61		
THYRO1000395	0.8	10.44	2.26	1.97	2.07	2.02		
THYRO1000400	0.57	8.1	2.82	2.35	2.96	2.52		
THYRO1000401	0.86	1.94	2.5	1.87	1.16	1.57		
THYRO1000407	1.97	2.3	1.36	1.37	1.58	0.55		
THYRO1000420	1.8	2.67	4.46	3.52	3.53	3.39		
THYRO1000438	1.78	4.37	3.26	2.94	3.33	3.15		
THYRO1000452	2.62	7.99	6.45	3.71	5.75	4.38		
THYRO1000455	0.32	6.67	2.31	0.25	0.97	0.87		
THYRO1000471	0.99	8.03	2.05	1.11	2.08	1.02		
THYRO1000481	1.33	6.23	4.68	3.79	3.45	4.55		
THYRO1000484	1.2	1.42	2.41	2.35	3	2.21		
THYRO1000488	1.18	2.64	2.44	1.49	2.02	1.7		
THYRO1000501	1.12	4.01	2.78	3	1.92	1.82		
THYRO1000502	0.34	3.7	1.69	1.79	1.44	1.2		
THYRO1000505	0.13	4.64	1.19	1.14	1.02	0.6		
THYRO1000535	11.1	20.54	39.24	54.13	69.59	62.96	*	+
THYRO1000556	1.89	6.36	4.13	3.77	5.17	3.69		
THYRO1000558	0.25	2.82	1.12	1.16	0.81	0.61		
THYRO1000569	2.88	4.12	6.05	5.78	4.46	4.88		
THYRO1000570	2.31	3.28	8.46	8.53	6.04	3.49		
THYRO1000572	0.43	2.04	1.11	0.17	0.97	-0.42		
THYRO1000573	0.69	4.02	1.73	2.02	2.2	1.78		

(1208)

特開2002-191363

2413

2414

THYRO1000577	1.06	5	1.34	0.96	1.22	0.71
THYRO1000580	0.79	3.72	3.01	2.82	2.2	1.79
THYRO1000584	2.18	6.88	8.8	7.57	6.61	7.58
THYRO1000585	4.83	9.37	9.83	5.76	6.27	9.52
THYRO1000596	0.22	0.93	1.19	0.44	1.36	0.21
THYRO1000602	2.08	2.95	4.01	4.05	4.65	4.97
THYRO1000605	0.37	3.01	0.98	2.13	2.14	1.56
THYRO1000615	1.02	3.62	1.24	1.55	1.36	1.29
THYRO1000625	0.71	5.48	2.28	2.46	2.9	1.78
THYRO1000636	3.67	5.65	6.9	6.53	7.84	6.67
THYRO1000637	0.91	3.96	1.71	1.18	2.03	1.54
THYRO1000641	0.38	4.19	2.49	1.36	1.67	1.64
THYRO1000657	2.99	3.69	5.42	7.67	12.28	3.86
THYRO1000658	2.68	3.62	5.39	5.4	5.55	6.09
THYRO1000662	1.1	3.19	2.09	2.42	2.69	1.66
THYRO1000666	0.57	3.19	2.28	1.63	1.48	1.43
THYRO1000676	1.37	4.53	2.01	1.75	1.83	1.56
THYRO1000678	0.52	5.86	0.99	1.29	1.4	0.53
THYRO1000684	0.95	4.98	2.94	1.92	2.65	1.47
THYRO1000694	2.08	6.64	4.65	2.8	2.48	3.59
THYRO1000699	2.98	2.14	5.55	4.86	7.08	7.12
THYRO1000712	1.88	4.25	5.9	6.25	6.75	7.78
THYRO1000715	5.74	5.67	27.37	21.74	28.63	16.99
THYRO1000716	0.92	3.26	3.2	1.88	1.78	1.35
THYRO1000717	1.58	5	4.36	2.98	4.63	1.91
THYRO1000723	0.6	4.54	1.6	0.55	1.06	0.85
THYRO1000734	-0.01	4.81	1.89	1.49	1.73	1.07
THYRO1000748	0.98	5.51	5.23	2.35	3.85	3.18
THYRO1000755	1.74	3.26	4.32	4.33	3.47	4.38
THYRO1000756	2.79	4.24	3.24	3.46	4.2	3.41
THYRO1000776	0.48	2.17	3.02	3.36	3.99	3.34
THYRO1000777	1.81	3.39	4.54	4.99	2.05	2.37
THYRO1000779	1.45	3.55	0.88	0.18	1.01	-0.26
THYRO1000782	3.92	10.13	12.52	10.76	15.05	14.05
THYRO1000783	0.12	5.51	1.2	1.11	1.41	0.92
THYRO1000786	6.65	9.54	19.71	15.74	7.92	13.7
THYRO1000787	0.23	1.88	1.67	1.31	1.54	0.78
THYRO1000792	1.51	3.13	2.29	3.09	3.13	2.11
THYRO1000793	0.11	3.13	0.84	1.51	1.86	1.16
THYRO1000795	1.23	6.03	3.54	2.76	3.1	3.05
THYRO1000796	0.6	7.73	2.44	2.26	2.95	1.66
THYRO1000798	1.89	5.82	2.51	2.59	3.57	3.53
THYRO1000800	9.26	17.2	24.74	17.74	20.68	21.06
THYRO1000805	0.49	3.04	1.08	0.72	2.66	1.38
THYRO1000815	2.54	3.49	9.48	7.61	5.47	7.87
THYRO1000829	5.55	7.83	10.57	3.78	8.32	10.01
THYRO1000835	0.96	3.2	1.93	1.07	2.36	1.8
THYRO1000843	1.09	11.48	3.56	3.69	4.41	3.62
THYRO1000846	0.76	5.71	1.32	2.67	1.62	1.26
THYRO1000852	1.59	6.02	5.63	2.8	4.7	3.32

(1209)

特開 2002-191363

2415

2416

THYRO1000855	3.14	5.02	6.63	9.03	15.1	10.07
THYRO1000865	1.86	4.3	11.97	10.01	11.47	8.95
THYRO1000866	7.47	6.29	12.66	4.49	7.87	6.01
THYRO1000881	5.62	7.3	10.93	15.65	26.64	29.58
THYRO1000894	0.33	3.95	1.36	1.75	1.48	1
THYRO1000895	0.58	4.43	1.42	1.62	1.46	0.82
THYRO1000916	1.22	5.49	3.43	2.43	3.13	2.29
THYRO1000917	16.19	25.26	34.11	30.37	37.42	35.89
THYRO1000926	0.78	3.13	1.27	1.76	1.57	0.82
THYRO1000934	0.08	3.1	1.34	0.43	1.38	1.46
THYRO1000951	0.52	2.46	1.26	2.33	2.11	1.9
THYRO1000952	2.25	3.81	6.01	2.38	2.53	2.24
THYRO1000956	0.06	2.55	1.81	1.16	1.5	0.87
THYRO1000960	0.5	6.72	2.89	1.85	2.79	1.48
THYRO1000961	1.67	7.77	3.56	4.73	5.26	4.64
THYRO1000964	0.42	11.59	0.76	1	1.27	1.06
THYRO1000971	1.82	9.9	3.56	3.29	3.33	2.19
THYRO1000974	2.87	8.83	7.53	9.87	11.79	8.71
THYRO1000975	1.5	2.19	3.8	4.02	3.68	3.68
THYRO1000983	6.42	8.31	11.63	12.67	8.49	7.12
THYRO1000984	2.4	2.83	3.03	3.29	2.98	3.26
THYRO1000988	1.36	4.14	3.23	3.48	3.68	2.67
THYRO1000991	1.22	4.71	2.05	1.76	2.22	3.2
THYRO1000999	0.87	9.64	3.26	1.96	3.14	2.26
THYRO1001003	2.97	8.43	4.42	3.1	4.52	3.56
THYRO1001015	0.6	6.29	2.04	2.22	1.79	1.66
THYRO1001016	1.73	2.26	3.34	2.06	1.85	1.24
THYRO1001022	0.9	1.86	0.86	1.68	1.25	1.41
THYRO1001031	4.65	3.97	4.55	5.03	7.03	6.16
THYRO1001033	1.18	3.34	2.46	2.86	3.45	2.02
THYRO1001062	1.21	5.4	4.14	2.9	4.31	2.62
THYRO1001063	0.5	8.74	2.38	2.37	2.09	2.84
THYRO1001071	0.12	7.45	0.88	1.33	0.68	0.76
THYRO1001080	2.56	6.75	5.11	4.96	4.31	4.78
THYRO1001093	0.77	1.63	3.24	5.11	1.74	1.5
THYRO1001100	0.52	1.89	2.05	1.89	1.21	0.78
THYRO1001102	2.61	3.6	5.7	4.4	4.95	6.93
THYRO1001104	3.67	6.54	6.55	8.77	8.01	11.18
THYRO1001109	1.81	6.02	2.68	3.06	2.58	1.99
THYRO1001113	11.41	17.42	32	21.81	26.65	18.72
THYRO1001120	1.65	6.22	5.27	4.78	5.8	3.72
THYRO1001121	1.57	4.28	4.19	3.92	2.72	3
THYRO1001128	1.64	2.77	5.86	3.52	3.19	5.09
THYRO1001133	1.14	3.02	7.23	6.54	4.54	4.12
THYRO1001134	2.97	4.78	1.63	3.14	2.83	1.38
THYRO1001142	0.3	2.69	0.63	1.3	1.71	0.22
THYRO1001173	8.37	12.87	7.72	11.14	9.92	10.62
THYRO1001175	3.26	6.63	5.51	3.46	4.62	3.52
THYRO1001177	1.36	5.85	5.93	4.66	7.27	7.97
THYRO1001189	2.74	6.93	11.42	7.84	7.27	9.94

(1210)

特開 2002-191363

2417

2418

THYRO1001194	1.05	2.62	4.96	4.89	4.57	2.31
THYRO1001204	2.17	3.58	4.27	4.03	4.74	4
THYRO1001205	5.76	10.65	20.23	18.54	19.57	20.9
THYRO1001213	1.21	4.69	4.44	3.12	3.21	2.33
THYRO1001224	3.59	8.25	6.37	9.92	12.69	10.55
THYRO1001237	2.82	6.25	4.99	3.61	4.53	4.46
THYRO1001242	9.74	11.65	19.04	20.02	19.98	20.46
THYRO1001258	2.08	5.45	3.58	3.33	2.05	2.66
THYRO1001262	0.86	2.64	3.38	2.36	3.61	2.69
THYRO1001266	0.15	2.39	1.02	0.97	1.64	0.66
THYRO1001271	1.85	4.12	4.12	2.46	2.77	2.97
THYRO1001287	7.3	8.3	39.26	30.14	43.68	26.2
THYRO1001290	0.38	3.25	1.14	1.15	1.35	0.36
THYRO1001291	0.96	7.17	4.38	4.31	4.97	3.5
THYRO1001297	3.05	8.04	6.14	6.85	7.47	9.18
THYRO1001302	1.72	5.59	5.17	3.8	3.71	3.5
THYRO1001313	1.61	2.33	2.91	2.91	2.62	1.48
THYRO1001320	1.76	2.52	5.31	5.07	5.74	4.83
THYRO1001321	2.25	2.65	4.3	2.48	4.23	4.23
THYRO1001322	1.34	3.93	3.34	1.75	2.67	2.01
THYRO1001327	1.29	6.01	4.18	1.4	3.89	2.49
THYRO1001336	1.89	6.84	6.72	4.62	4.43	4.18
THYRO1001347	0.43	4.12	3.35	1.85	2.81	0.65
THYRO1001358	2.57	5.74	4.52	4.3	5.75	5.1
THYRO1001363	0.8	2.15	1.52	2.09	2.24	2.28
THYRO1001365	0.86	3	1.6	2.19	2.6	1.96
THYRO1001374	1.85	4.45	12.86	9.4	13.01	6.21
THYRO1001401	1.76	5.33	4.89	5.39	7.86	6.29
THYRO1001403	1.26	5.15	3.22	3.22	4.42	3.94
THYRO1001405	6.99	12.5	10.86	6.69	8.56	10.63
THYRO1001406	15.73	14.87	27.69	21.36	22.77	21.36
THYRO1001411	4.49	5.46	10.08	8.93	12.44	8.12
THYRO1001420	11.55	15.25	47.52	42.01	44.49	49.87
THYRO1001426	3.42	5.56	8.83	9.32	12.77	11.18
THYRO1001430	6.97	6.54	10.84	11.13	11.7	13.81
THYRO1001434	0.68	5.19	2.11	2.08	4.23	1.73
THYRO1001456	1.74	6.05	2.63	2.89	2.66	2.4
THYRO1001457	1.71	4.72	2.04	2.95	4.7	2.67
THYRO1001458	0.95	5.44	6.11	6.13	9.17	7.23
THYRO1001459	4.54	5.07	9.42	7.18	9.87	14.21
THYRO1001471	0.91	2.07	1.93	2.36	2.91	1.64
THYRO1001478	0.58	3.09	1.34	0.95	2.61	2.75
THYRO1001480	5.4	10.53	13.62	14.79	15.94	15.57
THYRO1001481	2.95	8.64	7.24	4.91	7.76	7.13
THYRO1001487	1.36	5.51	3.52	4.1	2.67	3.48
THYRO1001495	2.06	5.57	5.43	5.06	9.4	7.55
THYRO1001498	5.39	8.08	13.42	11.59	16.38	19.6
THYRO1001510	1.67	2.88	1.59	1.65	2.57	3.35
THYRO1001512	26.7	26.97	110.28	56.03	90.95	68.54
THYRO1001519	5.92	7.77	11.05	8.91	5.75	10.38

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(1211)

特開 2002-191363

2419

2420

THYRO1001522	2.02	4.7	5.79	4.15	4.69	5.51		
THYRO1001523	1.83	4.92	3.73	4.38	4.07	4.26		
THYRO1001526	26.21	28.22	44.73	34.28	47.64	46.52		
THYRO1001529	1.64	5.27	2.8	2.32	2.42	2.98		
THYRO1001534	1.41	4.01	4.96	5.1	5.9	4.54		
THYRO1001537	7.4	5.17	12.33	5.97	7.42	7.23		
THYRO1001541	2.14	3.88	8.27	7.76	8.7	6.73		
THYRO1001545	1.26	3.84	2.9	4.95	3.57	3.16		
THYRO1001559	4.52	6.34	8.04	9.06	10.5	10.54	*	+
THYRO1001563	9.49	14.06	15.89	10	15.49	22.09		
THYRO1001570	2.01	8.2	3.85	4.25	5.17	3.41		
THYRO1001573	1.15	5.77	2.22	1.47	2.87	2.67		
THYRO1001584	2.47	8.54	8.38	5.14	7.81	7.29		
THYRO1001593	4.27	5.67	11.17	9.5	10.93	9.52		
THYRO1001595	3.14	4.53	7.06	5.97	6.35	7.29		
THYRO1001596	4.71	5.48	7.44	6.45	5.86	2.51		
THYRO1001602	1.49	3.26	3.52	4.95	5.22	3.41		
THYRO1001605	1.58	4.48	3.22	3.2	3.43	2.42		
THYRO1001608	1.87	9.45	5.1	5.04	8.23	4.7		
THYRO1001617	6.06	13.68	11.47	9.75	10.87	9.61		
THYRO1001634	1.87	9.08	3.46	2.93	5.59	2.05		
THYRO1001637	3.51	3.13	9.65	8.72	7.94	9.07		
THYRO1001641	2.57	3.73	5.09	4.03	3.08	2.94		
THYRO1001656	1.59	2.94	4.16	2.82	5.36	2.33		
THYRO1001658	22.34	29.19	40.11	34.98	33.16	42.01		
THYRO1001661	1.4	5.83	2.31	2.93	3.31	2.05		
THYRO1001671	0.67	7.36	2.68	1.89	1.34	1.8		
THYRO1001672	1.1	9.24	2.1	1.14	1.52	1.66		
THYRO1001673	1.59	7.6	3.49	2.86	4.74	2.16		
THYRO1001677	1.6	2.27	3.87	3.03	3.54	3.36		
THYRO1001683	12.71	17.66	29.06	24.4	15.4	16.72		
THYRO1001700	1.39	2.52	2.67	2.09	1.58	1.37		
THYRO1001702	11.83	15.98	16.19	15.63	14.35	14.29		
THYRO1001703	1.63	6.74	4.25	4.72	3.27	4.21		
THYRO1001706	1.7	6.47	3.01	2.96	5.6	3.53		
THYRO1001721	1.84	5.66	3.2	2.73	6.37	2.77		
THYRO1001725	5.3	6.55	9.69	8.97	8.65	8.29		
THYRO1001730	17.72	20.4	40.1	30.61	26.56	34.8		
THYRO1001738	1.35	3.18	4.65	3.52	2.82	1.78		
THYRO1001743	0.19	2.13	1.85	1.8	1.64	1.06		
THYRO1001745	0.47	2.88	1.55	1.05	1.2	1.27		
THYRO1001746	1.9	6.25	4.04	6.12	4.01	3.88		
THYRO1001770	15.49	20.38	35.39	41.65	44.42	40.17	*	+
THYRO1001772	1.12	4.88	3.64	4.78	4.24	3.06		
THYRO1001778	3.89	6.68	9.89	14.67	13.47	14.25	*	+
THYRO1001793	3.85	3.77	9.43	10.3	10.42	4.92		
THYRO1001796	1.35	2.28	2.28	3.45	4.22	3.24	*	+
THYRO1001800	1.82	2.99	2.75	4.17	5.12	2.09		
THYRO1001803	3.42	6.03	5.21	4.31	4.14	3.42		
THYRO1001809	1.6	4.26	3.4	5.9	4.23	3.39		

(1212)

特開2002-191363

2421

2422

THYRO1001817	8.69	18.33	24.88	22.11	22.77	25.27		
THYRO1001819	4.68	8.46	9.01	7.84	10.46	6.77		
THYRO1001828	21.89	24.53	104.32	106.68	121.95	68.55		
THYRO1001854	6.67	6.12	12.45	13.43	15.86	13.31		
THYRO1001895	0.85	1.31	2.52	3.54	4.31	1.28		
THYRO1001907	2.16	3.08	3.37	3.9	4.53	2.74		
TRACH1000006	2.51	5.87	6.63	4.34	3.46	3.22		
TRACH1000013	1.53	4.65	3.68	2.55	3.33	2.65		
TRACH1000074	2.65	6.75	6.09	7.5	7.26	4.77		
TRACH1000095	0.28	5.66	2.23	2.46	1.48	1.11		
TRACH1000102	2.42	6.66	4.09	5.79	5.04	3.65		
TRACH1000108	1.1	1.01	2.05	1.75	2.49	1.09		
TRACH1000126	0.96	1.75	2.71	1.82	3.79	2.54		
TRACH1000146	1.3	2.67	2.31	3.02	5.18	3.75		
TRACH1000160	0.61	4.06	1.5	1.47	1.76	0.72		
TRACH1000184	4.45	7.16	10.16	7.47	8.73	5.69		
VESEN1000004	0.69	5.55	3.19	2.56	2.95	2.02		
VESEN1000007	0.93	5.32	2.94	2.38	3.45	2.94		
VESEN1000013	5.96	10.11	16.78	10.76	11.25	13.88		
VESEN1000028	5.2	7.5	9.88	13.18	11.71	14.08	*	+
VESEN1000059	1.55	2.88	2.1	3.38	2.82	2.27		
VESEN1000100	1.96	3.22	3.35	3.49	4.58	3.59		
VESEN1000107	0.88	4.84	2.88	3.12	2.9	2.48		
VESEN1000117	1.63	6.43	2.46	2.16	2.7	1.79		
VESEN1000122	1.52	5.34	1.24	4.79	4.51	4.5		
VESEN1000137	0.76	5.47	1.92	1.75	3.33	1.65		
VESEN1000195	7.79	7.93	11.67	8.42	7.51	10.27		
VESEN1000215	1.48	3.03	2.06	2.67	3.84	1.87		
VESEN1000279	8.71	11.32	18.49	22.93	23.38	34.68	*	+
VESEN1000363	3.52	6.07	9.99	7.2	9.06	4.59		
VESEN1000388	2.55	6.48	3.31	4.17	3.75	6.7		
VESEN1000394	0.44	7.11	2.33	2.37	2.55	2.36		
VESEN1000410	1.11	5	1.78	2.36	2.71	3.69		
VESEN1000411	2.37	4.95	5.08	6.76	7.55	9	*	+
VESEN1000415	1.54	2.64	4.03	5.57	3.92	5.29		
VESEN1000440	7	5.53	7.81	3.79	9.4	12.22		
VESEN1000452	1.22	3.65	2.33	2.91	3.97	4.11		
VESEN1000539	191.54	185.28	334.6	389.84	403.89	547.31	*	+
VESEN1000554	0.67	6.47	1.43	1.47	2.55	1.58		
VESEN1000557	4.22	7.94	7.73	6.55	9.07	10		
VESEN1000575	7.49	9.75	16.33	11.95	11.73	14.8		
VESEN1000585	1.69	4.49	3.37	2.53	2.93	3.08		
VESEN1000592	1.58	2.31	1.58	2.02	1.83	1.46		
VESEN1000658	1.96	3.56	4.45	5.86	3.91	4.91		
VESEN1000669	8.43	10.02	16.35	15.06	14.17	15.51		
VESEN1000743	0.97	3.3	2.52	1.99	3.37	3.21		
VESEN1000752	37.43	51.51	72.35	49.32	57.03	57.96		
VESEN1000761	13.48	18.17	19.37	24.6	33.21	31.28	*	+
VESEN2000039	10.45	15.98	15.56	13.56	18.3	20.1		
VESEN2000102	0.4	3.99	1.6	1.21	1.51	1.61		

(1213)

特開2002-191363

2423

2424

VESEN2000164	2.45	3.52	4.8	5.55	4.36	3.66		
VESEN2000175	0.57	2.64	1.94	3.03	2.05	2.59		
VESEN2000186	3.77	5.53	6.53	6.68	3.87	2.47		
VESEN2000199	8.94	13.26	21.75	19.58	24.45	24.12		
VESEN2000200	0.5	4.97	2.78	3.03	3.1	1.6		
VESEN2000204	0.48	12.7	1.02	0.98	1.2	0.33		
VESEN2000218	6.66	20.26	19.48	21.37	20.75	18.86		
VESEN2000230	0.84	7.4	1.45	2.74	1.7	2.41		
VESEN2000272	2.29	4	8.92	7	8.31	5.88		
VESEN2000299	1.99	2.97	3.2	3.3	3.77	3.31		
VESEN2000323	4.51	8.12	8.37	8.91	9.5	9.36		
VESEN2000327	3.16	5.8	4.42	3.62	7.66	5.07		
VESEN2000328	5.44	8.02	11.88	8.73	15.35	14.36		
VESEN2000330	6.39	15.42	14.7	14.59	27.73	18.86		
VESEN2000336	0.82	8.97	2.54	3.46	3.83	2.88		
VESEN2000354	1.56	8.24	2.48	1.53	2.71	2.1		
VESEN2000378	7.17	8.87	14.57	13.23	11.14	10.3		
VESEN2000379	19.87	23.02	44.55	49.13	42.81	32.61		
VESEN2000397	0.72	2.38	1.24	1.36	2.06	1.54		
VESEN2000416	2.83	3.88	4.41	5.74	5.31	5.71	*	+
VESEN2000420	1.08	3.58	1.94	0.95	1.21	1.13		
VESEN2000430	0.51	6.68	1.53	2.06	1.79	1.8		
VESEN2000448	0.51	6.87	1.73	2.12	2.69	1.5		
VESEN2000449	2.43	8.07	6.59	8.3	11.31	8.21		
VESEN2000456	0.74	0.87	2.11	1.54	1.22	0.87		
VESEN2000562	4.07	3.42	17.42	13.67	22.82	14.47		
VESEN2000573	0.18	1.75	1.75	1.04	1.21	1.13		
VESEN2000604	1.73	3.44	2.24	2.04	1.67	2.15		
VESEN2000614	4.16	9.02	14.64	13.54	16.27	12.27		
VESEN2000638	0.48	5.92	1.98	1.33	1.91	1.68		
VESEN2000641	0.83	3.69	1.34	1.95	2.21	1.43		
VESEN2000645	2.18	5.29	5.38	5.91	5.73	5.5		
Y79AA1000013	2.57	2.7	4.33	3.45	3.8	3.94		
Y79AA1000030	1.79	4.06	4.52	3.24	3.85	2.47		
Y79AA1000033	2.87	6	8.4	8.37	10.17	5.83		
Y79AA1000037	1.38	3.36	5.71	4.84	6.82	4.49		
Y79AA1000041	1.05	5.16	3.79	4.73	3.65	2.06		
Y79AA1000059	1.69	5	4.09	3.88	4.51	2.82		
Y79AA1000065	24.06	28.99	52.25	82.48	101.48	98.73	**	+
Y79AA1000081	39.47	49.78	73.62	113.19	114.49	98.22	**	+
Y79AA1000127	4.08	4.21	5.8	8.42	10.03	7.39	*	+
Y79AA1000130	2.24	2.48	5.76	6.61	8.1	8.03	*	+
Y79AA1000131	507.64	569.21	946.04	769.75	725.35	342.07		
Y79AA1000134	1.99	4.93	4.21	5.63	4.75	5.38		
Y79AA1000143	3.58	8.79	4.83	9.98	10.98	11.04	*	+
Y79AA1000144	4.63	10.79	10.59	11.02	11.62	11		
Y79AA1000150	18.39	22.18	84.69	93.5	117.62	78.89		
Y79AA1000153	183.67	191.4	436.64	423.46	442.52	386.45		
Y79AA1000166	2.13	2.25	4.15	3.52	4.97	3.6		
Y79AA1000179	2.58	3.76	4.2	6.85	7.58	3.89		

(1214)

特開2002-191363

2425

2426

Y79AA1000181	1.96	3.92	4.2	4.82	5.66	3.79		
Y79AA1000202	22.93	24.47	55.57	91.68	86.86	83.22	**	+
Y79AA1000207	5.22	7.51	9.82	14.95	16.24	12.62	*	+
Y79AA1000214	14.94	22.18	33.76	50.43	60.78	41.96	*	+
Y79AA1000222	11.8	14.89	21.69	49.21	58.01	68.86	**	+
Y79AA1000226	11.04	14.94	34.41	22.86	33.54	30.03		
Y79AA1000227	5.95	4.52	7.25	8.98	9.51	9.62	*	+
Y79AA1000230	1.09	1.49	2.02	2.07	2.88	2.59		
Y79AA1000231	5.99	9.04	15.81	14.63	23.77	17.1		
Y79AA1000239	15.47	20.55	25.65	18.95	24.01	22.11		
Y79AA1000258	2.64	5.17	6.72	5.87	4.95	5.84		
Y79AA1000268	2.65	5.48	5.09	4.33	5.76	4.01		
Y79AA1000269	4.32	7.88	7.86	8	6.86	8.24		
Y79AA1000270	5.28	8.35	11.58	13.17	17.58	16.23	*	+
Y79AA1000280	1.74	4.17	4.9	5.29	3.1	5.27		
Y79AA1000285	3.44	4.21	5.91	4.01	6.86	5.22		
Y79AA1000295	0.75	3.06	4.85	4.32	4.23	4.45		
Y79AA1000307	2.88	3.91	5.06	9.35	7.58	11.25	*	+
Y79AA1000313	3.11	9.02	9.85	10.61	12.84	13.36		
Y79AA1000314	4.23	10.74	9.19	6.93	6.53	7.51		
Y79AA1000328	4.65	10.05	2.64	7.73	9.28	8.68		
Y79AA1000334	1.43	4.22	3.55	2.68	2.81	3.46		
Y79AA1000342	10.65	10.05	26.47	23.7	19.13	28.35		
Y79AA1000346	7.61	8.17	7.9	20.1	22.06	20.18	**	+
Y79AA1000347	6.94	7.96	12.42	18.78	16.47	18.48	**	+
Y79AA1000349	6.93	9.63	12.67	12.31	11.96	14.75		
Y79AA1000355	3.17	8.28	8.94	8.84	13.55	8.77		
Y79AA1000368	5.24	8.39	24.43	22.48	35.67	22.55		
Y79AA1000388	22.9	32.66	62.37	94.17	128.35	109.08	**	+
Y79AA1000392	3.02	6.81	3.76	3.42	2.73	3.78		
Y79AA1000405	3.98	5.97	8.25	7.14	8.17	9.32		
Y79AA1000410	6.01	7.87	15.72	13.79	17.05	14.95		
Y79AA1000420	1.54	4.78	3.13	3.32	3.95	5.1		
Y79AA1000423	1.38	7.08	5.59	5.22	6.04	10.27		
Y79AA1000426	3.61	9.44	8.66	4.24	4.43	5.22		
Y79AA1000432	0.8	4.79	2.16	1.91	2.01	2.34		
Y79AA1000453	23.94	30.67	47.79	39.74	50.65	58.24		
Y79AA1000465	4.12	6.02	6.65	4.77	4.14	7.69		
Y79AA1000469	11.59	9.61	18.04	13.82	16.21	17.18		
Y79AA1000480	1.24	4.37	2.78	3.33	3.57	2.79		
Y79AA1000502	5.31	7.97	12.58	10.49	11.35	15.26		
Y79AA1000521	1.24	4.4	4.13	2.51	3.61	2.7		
Y79AA1000534	3.22	8.13	8.92	11.97	14.41	13.46	*	+
Y79AA1000538	3.58	6.95	8.79	9.52	12.12	8.41		
Y79AA1000539	12.76	14.96	53.11	42.61	68.56	50.97		
Y79AA1000540	1.32	3.59	1.61	2.54	2.97	3.21		
Y79AA1000560	160.46	140.99	339.33	380.8	313.21	220.43		
Y79AA1000574	1	2.92	1.65	1.98	2.04	1.59		
Y79AA1000584	2.07	4.55	4.97	4.62	5.39	4.04		
Y79AA1000589	10.74	13.67	81.43	59.09	95.35	68.5		

(1215)

特開 2002-191363

2427

2428

Y79AA1000598	1.43	7.64	2.17	1.85	3.88	3.18		
Y79AA1000600	2.7	10.02	7.93	13.64	15.64	12.84	*	+
Y79AA1000609	1.18	5.16	1.44	2.28	2.77	1.55		
Y79AA1000618	1.85	10.59	5.76	7.4	9.5	9.37		
Y79AA1000627	1.91	3.93	4.57	3.27	3.02	2.43		
Y79AA1000636	5.16	5.7	9.9	15.57	11.52	5.38		
Y79AA1000649	9.45	10.97	12.73	18.7	11.56	20.54		
Y79AA1000656	15.32	20.21	96.75	80.17	115.97	82.61		
Y79AA1000673	1.02	5.86	2.14	1.39	3.13	2.4		
Y79AA1000674	11.88	21.96	78.28	59.4	98.22	62.67		
Y79AA1000678	2.48	8.91	3.88	3.01	4.15	2.45		
Y79AA1000682	17.99	53.99	93.7	102.53	110.87	118.22		
Y79AA1000683	1.87	2.66	3.21	4.27	2.59	2.17		
Y79AA1000697	21.76	27.52	43.01	21.93	24.76	27.31		
Y79AA1000700	5.07	7.1	7.08	7.51	6.93	9.97		
Y79AA1000702	5.13	14.57	13.31	41.48	56.57	63.15	**	+
Y79AA1000704	1.34	5.24	1.14	1.5	2.1	1.18		
Y79AA1000705	1.86	9.7	6.24	6.06	7.98	6.45		
Y79AA1000717	6.18	12.39	9.42	9.73	11.1	8.7		
Y79AA1000722	5.61	8.6	8.65	26.26	34.81	34.31	**	+
Y79AA1000724	6.42	9.77	18.55	26.57	21.7	11.95		
Y79AA1000726	0.77	1.24	0.83	1.46	2.01	0.71		
Y79AA1000734	2.05	4.46	4.92	3.57	3.3	2.94		
Y79AA1000748	0.88	4.38	1.77	2.14	2.99	1.56		
Y79AA1000750	4.17	8.47	20.58	18.11	20.02	18.62		
Y79AA1000752	1.25	5.13	2.33	2.23	3.38	3.38		
Y79AA1000774	2.11	6.23	4.24	4.28	5.79	4.21		
Y79AA1000776	1.2	4.37	1.9	2.13	1.83	2.11		
Y79AA1000777	4.36	5.84	9.63	10.05	6.99	6.01		
Y79AA1000778	1.72	3.77	3.79	3.87	4.19	1.44		
Y79AA1000782	2.08	4.18	3.72	3.53	2.89	2.96		
Y79AA1000784	7.04	10.01	7.78	13.87	15.58	14.26	**	+
Y79AA1000794	0.61	5.21	1.88	2.92	1.69	1.23		
Y79AA1000800	1.59	5.44	3.82	3.38	2.97	3.39		
Y79AA1000802	0.64	4.18	1.15	2.52	1.77	2.1		
Y79AA1000805	2.29	4.03	2.63	2.43	2.11	2.6		
Y79AA1000814	2.73	3.9	4.14	4.98	7.09	6.76	*	+
Y79AA1000823	7.91	9.99	12.07	12.02	12.42	6.56		
Y79AA1000824	0.98	2.47	1.84	2.75	2.26	0.79		
Y79AA1000827	1.6	4.02	7.27	6.71	8.91	6.14		
Y79AA1000831	7.04	10.49	17.32	26.61	30.43	27.82	**	+
Y79AA1000833	62.14	67.46	191.76	270.42	308.16	248.21	*	+
Y79AA1000850	1.69	5.68	2.72	4.92	4.59	4.21		
Y79AA1000856	3.49	6.78	6.31	9.28	6.01	8.51		
Y79AA1000862	2.22	2.76	2.44	3.08	2.87	2.59		
Y79AA1000876	7.46	10.04	17.91	27.36	27.61	25.46	**	+
Y79AA1000888	4.59	5.07	28.1	24.51	38.78	22.48		
Y79AA1000902	4.65	5.74	8.44	12.18	12.32	7.68		
Y79AA1000935	3.53	5.99	6.69	8.28	10.07	9.18	*	+
Y79AA1000959	0.74	6.29	4.35	6.71	5.77	6.07		

(1216)

特開 2002-191363

2429

2430

Y79AA1000962	1.22	4.45	3.18	2.9	2.41	1.79		
Y79AA1000963	18.6	26.86	35.93	31.61	42.17	49.13		
Y79AA1000966	18.52	19.25	98.18	77.47	116.12	61.1		
Y79AA1000967	8.62	8.82	33.82	34.47	40.36	29.3		
Y79AA1000968	3.32	5.67	6.89	8.86	9.4	7.96	*	+
Y79AA1000969	0.91	3.18	3.04	2.66	2.91	1.41		
Y79AA1000976	1.43	4.72	3.02	2.55	3.51	1.43		
Y79AA1000978	1.99	5.6	7.26	6.9	10.54	5.56		
Y79AA1000985	9.39	12.84	40.49	38.26	44.3	27.89		
Y79AA1000989	21.59	22.49	46.19	51.84	58.65	55.19	*	+
Y79AA1000991	22.11	22.21	110.42	72.46	96.96	82.23		
Y79AA1001013	59.2	62.64	140.9	174.85	214.13	201.9	*	+
Y79AA1001014	2.27	4.16	4.1	4.8	5.2	6.55		
Y79AA1001019	3.37	5.89	7.74	9.24	9.02	9.43	*	+
Y79AA1001020	5.37	7.82	9.43	12.31	11.11	10.86	*	+
Y79AA1001023	0.83	6.11	2.29	1.22	1.95	1.54		
Y79AA1001030	4.23	8.79	10.87	11.14	10.72	12.43		
Y79AA1001035	0.19	2.88	0.03	14.44	8.19	17.16	*	+
Y79AA1001041	1.78	2.46	2.36	2.93	2.45	2.78		
Y79AA1001043	11.65	12.62	15.22	8.64	12.01	14.71		
Y79AA1001048	1.1	4.78	3.73	4.05	4.52	4.21		
Y79AA1001056	4.56	7.82	11.04	8.27	7.11	9.94		
Y79AA1001061	1.53	7.79	5.28	6.13	7.46	6.66		
Y79AA1001062	2.62	6.14	5.02	4.44	6.01	4.67		
Y79AA1001068	3.46	6.39	7.29	6.61	8.69	7.05		
Y79AA1001073	8.19	13.08	17.46	24.14	22.1	29.81	*	+
Y79AA1001077	7.1	7.08	17.15	14.69	14.74	17.08		
Y79AA1001078	3.11	8.34	11.07	5.01	12.15	12.92		
Y79AA1001081	3.59	5.61	4.94	9.62	9.98	10.5	**	+
Y79AA1001088	27.75	38.61	69.33	93.1	88.97	113.04	*	+
Y79AA1001089	4.64	7.8	11.92	22.67	22.6	27.73	**	+
Y79AA1001090	1.38	4.15	2.2	3.58	2.83	2.35		
Y79AA1001105	3.7	5.23	15.81	12.52	22.1	13.35		
Y79AA1001142	8.53	13.38	15.85	14.28	11.42	22.32		
Y79AA1001145	2.22	4.68	5.13	4.97	6.26	5.87		
Y79AA1001162	2.27	2.91	1.62	1.62	4.56	4		
Y79AA1001167	0.86	2.76	2.38	1.12	2.35	0.77		
Y79AA1001176	0.57	3.33	1.14	2.02	1.68	0.88		
Y79AA1001177	1.21	5.5	2.22	2.35	3.01	1.99		
Y79AA1001179	6.81	8.66	16.73	22.82	22.64	20.07	*	+
Y79AA1001185	1.33	5.3	4.55	3.65	4.49	5.8		
Y79AA1001201	5.69	11.3	16.13	14.57	15.21	19.38		
Y79AA1001205	1.87	3.28	2.85	5.87	4.85	4.09	*	+
Y79AA1001211	1.64	4.75	6.93	4.83	4.36	4.15		
Y79AA1001212	3.55	6.93	15.91	13.74	15	11.65		
Y79AA1001216	52.59	51.46	93.73	76.52	97.53	109.55		
Y79AA1001228	6.1	11.21	9.34	8.99	12.19	10.24		
Y79AA1001233	0.68	11.46	2.39	0.92	1.66	1.09		
Y79AA1001236	4.46	12.86	9.25	11.4	10.66	13.08		
Y79AA1001239	4.62	13.93	9.94	11.53	12.15	12.94		

(1217)

特開2002-191363

2431

2432

Y79AA1001240	8.74	8.6	13.75	9.13	6.68	3.01		
Y79AA1001255	10.37	12.22	22.61	12.47	7.51	6.57		
Y79AA1001264	3.63	5.15	4.49	7.73	8.59	8.75	**	+
Y79AA1001272	10.81	13.63	17.47	21.56	20.67	21.32	*	+
Y79AA1001281	0.45	4.95	1.89	1.42	1.81	0.95		
Y79AA1001299	2.49	11.34	9.06	9.9	9.26	9.81		
Y79AA1001312	2.49	10.36	5.17	2.15	4.77	4.14		
Y79AA1001319	3.34	11.88	5.27	7.23	6.05	7.15		
Y79AA1001323	1.22	1.16	2.09	1.11	1.55	0.89		
Y79AA1001328	2.04	3.18	3.62	4.66	4.48	4.05	*	+
Y79AA1001343	154.19	151.55	345.05	304.88	394.54	265.65		
Y79AA1001351	0.81	2.77	1.67	0.03	1.26	0.96		
Y79AA1001364	1.65	6.07	4.03	3.39	4.43	3.6		
Y79AA1001367	2.16	7.41	2.93	3.09	5.34	3.19		
Y79AA1001384	0.5	5.14	1.98	0.73	1.15	0.94		
Y79AA1001391	0.59	3.73	2.88	1.35	1.65	1.2		
Y79AA1001394	3.12	4.66	12.92	10.94	9.56	10.94		
Y79AA1001402	2.77	3.7	5.95	5.65	5.09	4.14		
Y79AA1001410	0.82	2.78	2.33	2.06	2.31	2.25		
Y79AA1001414	2.76	7.5	7.59	11.08	10.73	10.06	*	+
Y79AA1001426	0.61	4.36	2.61	1.5	1.43	1.82		
Y79AA1001427	14.22	13.44	86.36	59.92	88.36	63.53		
Y79AA1001430	11.28	16.98	20.98	29.04	34.03	34.38	**	+
Y79AA1001439	16.22	21.53	33.42	45.02	43.74	43.1	*	+
Y79AA1001485	1.65	2.51	4.38	4.85	3.88	3.57		
Y79AA1001493	1.29	2.3	3.43	2.43	2.35	2.46		
Y79AA1001511	4.79	8.57	11.05	9.39	9.11	6.47		
Y79AA1001523	2.64	6.57	5.08	8.74	7.37	6.1		
Y79AA1001530	7.46	11.69	22	41.43	36.37	36.07	**	+
Y79AA1001532	5.12	7.35	6.69	10.49	14.82	12.9	*	+
Y79AA1001533	1.84	4.89	2.53	4.15	4	4.19		
Y79AA1001541	2.82	5.89	7.54	7.23	5.08	7.34		
Y79AA1001548	4.25	4.84	9.82	10.46	7.37	7.1		
Y79AA1001555	2	2.55	3.65	4.7	5.28	4.25	*	+
Y79AA1001562	7.76	10.11	17.15	14.07	16.16	10.83		
Y79AA1001581	2	5.05	4.47	5.1	7.01	3.54		
Y79AA1001585	3.18	7.38	10.96	9.72	10.93	8.05		
Y79AA1001592	2.61	7.38	5.97	8.15	8	7.02		
Y79AA1001594	0.76	4.73	3.85	1.96	3.24	1.73		
Y79AA1001603	56.74	70.81	153.14	131.56	112.16	107.66		
Y79AA1001613	3.74	3.52	14.81	13.12	15.36	10.66		
Y79AA1001630	0.71	2.36	1.73	1.14	2.64	0.67		
Y79AA1001647	1.96	3.57	2.47	4.14	4.32	2.65		
Y79AA1001664	4.67	8.39	11.43	8.96	10.01	8.73		
Y79AA1001665	1.39	6.4	3.73	4.67	4.71	3.75		
Y79AA1001679	8.92	15.94	20.71	20.53	26.67	25.35		
Y79AA1001692	1.87	5.55	3.95	3.99	3.95	3.51		
Y79AA1001696	1.97	6.49	2.77	1.83	2.63	2.55		
Y79AA1001705	6.09	6.44	10.39	7.62	7.92	7.85		
Y79AA1001711	16.17	12.34	29.74	13.73	23.83	21		

(1218)

特開2002-191363

2433

2434

Y79AA1001717	0.72	2.99	1.29	1.68	3.13	1.14		
Y79AA1001719	2.5	5.79	6.44	6.15	6.07	6.43		
Y79AA1001727	6.87	12.13	14.99	8.73	14.71	8.77		
Y79AA1001750	10.21	13.63	21.67	21.92	32.29	24.33		
Y79AA1001760	25.24	27.31	122.97	113.56	155.17	83.24		
Y79AA1001777	1.17	3.59	1.6	2.49	1.75	1.28		
Y79AA1001781	0.31	2.3	0.42	1.62	1.84	1.42		
Y79AA1001787	1	3.94	3.54	5.51	5.18	5.15		
Y79AA1001793	16.23	15.19	91.7	60.44	87.21	75.35		
Y79AA1001795	1.23	3.84	2.24	2.05	2.45	2.8		
Y79AA1001799	4.9	8.35	6.99	10.7	10.72	11.26	*	+
Y79AA1001800	2.25	8.3	10.1	8.49	10.51	9.9		
Y79AA1001801	1.77	6.44	4.87	7.67	5.91	7.33		
Y79AA1001803	0.74	2.15	1.72	1.85	1.68	1.17		
Y79AA1001805	6.05	6.88	12.15	10.21	8.32	11.15		
Y79AA1001807	3.37	6.33	12.56	11.76	17.8	16.79		
Y79AA1001827	1.7	4.41	3.12	3.43	3.6	2.52		
Y79AA1001846	1.82	6.52	5.51	6.52	5.09	5.07		
Y79AA1001848	0.86	5.57	2.75	5.98	4.78	5.14		
Y79AA1001853	1.38	6.16	2.76	3.24	4.56	3.63		
Y79AA1001863	0.86	5	2.53	2.93	3.83	3.85		
Y79AA1001866	2.29	3.2	5.81	3.53	3.98	3.46		
Y79AA1001874	0.12	2.18	-0.21	1.17	0.75	0.51		
Y79AA1001875	9.33	12.67	13.09	11.05	17.79	18.63		
Y79AA1001907	68.02	70.94	96.4	118.34	86.75	104.86		
Y79AA1001908	0.64	8.4	2.29	1.92	2.59	2.31		
Y79AA1001923	1.61	6.64	3.03	3.86	3.76	3.35		
Y79AA1001927	19.1	22.05	36.94	42.46	45.29	48.81	*	+
Y79AA1001930	4.07	6.65	8.07	7.92	12.42	12.21		
Y79AA1001932	2.84	4.41	8.47	11.51	9.1	8.57		
Y79AA1001933	2.14	3.27	3.69	4.34	7.99	6.65	*	+
Y79AA1001942	1.58	3.45	2.69	2.94	2.13	2.41		
Y79AA1001963	9.6	9.37	46.06	38.48	49.64	47.27		
Y79AA1001968	18.61	27.73	37.44	42.93	44.16	55.23	*	+
Y79AA1001983	1.81	6.35	4.28	3.97	5.86	4.47		
Y79AA1002000	2.55	5.35	4.55	4.42	3.21	2.83		
Y79AA1002004	13.1	18.87	27.47	23.72	29.45	40.93		
Y79AA1002008	2.51	3.73	3.79	4.54	2.19	2.85		
Y79AA1002012	1.37	3.22	2.81	3.22	2.29	2.87		
Y79AA1002017	1.34	2.53	2.46	3.51	3.07	2.82		
Y79AA1002022	2.99	4.94	5.93	7.32	7.51	6.01		
Y79AA1002027	2.02	6.33	2.67	2.69	4.03	4.09		
Y79AA1002050	2.53	8.12	4.22	6.68	6.91	5.11		
Y79AA1002058	13.69	21.8	70.12	59.07	70.89	55.33		
Y79AA1002060	6.38	13.17	20.54	17.14	21.12	24.23		
Y79AA1002062	4.33	5.18	8.15	8.54	6.66	5.51		
Y79AA1002065	33.54	39.97	72.6	49.46	30.04	41.81		
Y79AA1002067	10.11	11.64	17.24	16.25	9.42	8.13		
Y79AA1002069	0.97	1.79	0.54	1.55	1.44	0.66		
Y79AA1002070	10.16	33.47	44.36	52.16	71.15	73.35	*	+

(1219)

特開 2002-191363

2435

2436

Y79AA1002074	38.55	74.38	179.6	165.55	282.48	224.96		
Y79AA1002076	0.48	9.71	2.89	2.86	3.34	1.91		
Y79AA1002083	1.2	7.48	2.03	2.73	1.75	2.06		
Y79AA1002084	1.79	2.59	4.54	3.73	3.73	2.98		
Y79AA1002086	0.96	1.78	1.71	2.77	1.88	1.43		
Y79AA1002087	11.18	14.9	27.67	33.34	30.01	23.08		
Y79AA1002089	1.18	3.46	2.13	1.46	2.92	3.26		
Y79AA1002093	2.19	5.48	5.25	5.28	5.68	6.17		
Y79AA1002101	1.11	8.58	2.98	6.54	5.58	6.6		
Y79AA1002103	1.47	10.22	3.39	4.43	6.49	4.7		
Y79AA1002115	4.34	9.78	7.37	7.45	7.03	6.95		
Y79AA1002121	1.55	2.16	2.18	1.67	2.55	2.31		
Y79AA1002125	6.67	7.08	9.29	8.81	6.4	7.6		
Y79AA1002129	1.64	6.23	7.84	5.41	2.2	4.93		
Y79AA1002131	0.9	3.25	0.78	0.77	1.29	1.15		
Y79AA1002139	0.69	5.02	1.04	1.83	1.53	1.34		
Y79AA1002144	25.99	29.62	51.01	42.61	51.16	43.17		
Y79AA1002177	1.72	5.97	4.33	4.79	3.09	4.73		
Y79AA1002183	10.44	13.89	17.69	27.61	29.67	28.92	**	+
Y79AA1002202	3.97	7.15	8.34	18.27	10.12	17.85	*	+
Y79AA1002204	0.53	0.99	1.56	1.7	2.2	1.54		
Y79AA1002206	2.63	5.36	7.28	4.35	2.95	1.49		
Y79AA1002208	4.26	6.54	3.94	6.88	6.3	3.96		
Y79AA1002209	1.8	6.34	2.88	4.38	3.74	4.57		
Y79AA1002210	0.41	4.14	2.09	1.8	2.24	1.65		
Y79AA1002211	2.25	5.39	3.85	5.71	5.3	4.5		
Y79AA1002213	1.15	4.13	6.53	7.38	7.54	7.43		
Y79AA1002215	18.7	18.69	26.61	17.72	15.59	9.62		
Y79AA1002220	3.78	3.38	2.87	4.89	4.19	4.14	*	+
Y79AA1002226	8.54	8.9	9.75	13.06	14.2	4.41		
Y79AA1002229	1.35	3.88	3.38	2.95	2.79	2.67		
Y79AA1002234	3.24	6.82	3.94	4.29	7.74	6.88		
Y79AA1002235	5.6	7.55	6.43	8.78	9.74	9.47	*	+
Y79AA1002246	0.59	5.06	2.41	3.94	2.54	4.27		
Y79AA1002258	0.72	7.26	2.92	3.99	4.19	2.7		
Y79AA1002279	17.79	19.12	27.8	16.52	19.13	11.5		
Y79AA1002292	1.68	2.1	3.22	2.96	3.91	2.73		
Y79AA1002298	0.76	2.52	1.32	2.03	2.77	1.06		
Y79AA1002307	1.05	4.35	1.79	0.76	1.05	1.2		
Y79AA1002309	1.15	4.19	2.3	2.21	1.78	2.55		
Y79AA1002311	2.84	7.35	3.43	5.71	6.04	5.45		
Y79AA1002334	1.72	6.54	2.95	4.77	4.19	3.35		
Y79AA1002351	1.27	5.5	2.89	3.5	3.38	3.06		
Y79AA1002355	12.83	12.25	28.96	22.94	22.07	21.02		
Y79AA1002361	2.22	2.27	3.26	2.47	4.54	1.55		
Y79AA1002365	0.66	2.04	2.26	1.97	3.51	2.25		
Y79AA1002373	1.17	3.93	2.42	1.59	1.97	1.43		
Y79AA1002376	110.81	135.82	249.8	205.99	213.25	191.69		
Y79AA1002378	1.9	4.8	4.91	2.2	3.6	3		
Y79AA1002381	8.65	14.11	19.19	18.84	21.52	17.97		

2437							2438
Y79AA1002388	7.05	9.99	18.24	15.88	21.51	19.99	
Y79AA1002399	1.79	4.25	3.74	4.62	4.08	3.47	
Y79AA1002407	3.05	4.16	3.13	4.66	5.77	4.5	*
Y79AA1002413	3.21	6.78	8.05	6.46	8.32	6.87	+
Y79AA1002416	1.46	5	2.74	2.49	3.44	3.55	
Y79AA1002429	5.5	8.15	7.27	8	11.11	8.01	
Y79AA1002431	0.92	4.43	0.48	0.79	1.78	0.89	
Y79AA1002433	1.27	5.9	3.24	4.8	3.84	5.58	
Y79AA1002445	4.01	5.34	5.76	3.1	4.89	5.41	
Y79AA1002461	0.63	2.45	1.79	1.19	2.71	1.41	
Y79AA1002466	39.02	70.71	94.5	91.12	82.27	94.71	
Y79AA1002471	4.44	6.67	6.08	7.43	8.06	10.49	
Y79AA1002472	2.41	6.16	5.99	6.8	8.39	4.06	
Y79AA1002474	1.93	8.27	4.31	4.89	6.52	7.13	
Y79AA1002482	3.52	6.66	10.37	9.02	11.81	8.69	
Y79AA1002487	1.38	4.12	2.46	1.96	3.01	2.56	
Y79AA1002490	10.37	9.91	16.35	11.11	12.88	16.86	
Y79AA1002493	1.96	4.07	6.14	6.5	8.9	4.1	
ZRV6C1006278	0.61	4.08	2.22	1.81	1.58	2.11	

【0824】

【表353】紫外線未照射の皮膚由来繊維芽細胞、及び紫外線照射した皮膚由来繊維芽細胞の各cDNAの発現（表中には実施例で説明の無いクローンも含まれている）表中、UV_0hは紫外線未照射の皮膚由来繊維芽細胞を、UV_4h、UV_24hはそれぞれ紫外線照射4時間後、24時間後の皮膚由来繊維芽細胞を示し、これに続くExp1、Exp2、

およびExp3は、それぞれn=3の測定結果を表している。t test 0/4とt test 0/24は、それぞれ紫外線未照射と紫外線照射4時間後の有意差検定、紫外線未照射と紫外線照射24時間後の有意差検定の結果を示す。紫外線照射した皮膚由来繊維芽細胞における、4時間後と24時間後の発現量の増加を+で、減少を-で+/-として示す。有意差検定の結果を、*P<0.05、および**P<0.01で示した。

2439

2440

Class	UV 0h			UV 4h			UV 24h			t test	4h 24h
	Exp. 1	Exp. 2	Exp. 3	Exp. 1	Exp. 2	Exp. 3	Exp. 1	Exp. 2	Exp. 3		
GAPDH(Cr1)	0	1.29	0.1	0.9	0.08	1.18	1.49	0.47	0		
βactin(Cr2)	258.82	283.53	414.29	268.38	117.29	329.8	189.18	180.26	157.87	*	-
ADRG11000095	16.9	10.69	19.67	3.3	8.44	12.69	8	3.08	8	*	-
ADRG11000097	47.47	31.82	85.7	41.54	41.35	40.62	31.64	14.77	31.5		-
ADRG11000099	3.97	2.45	8.72	6.06	2.65	3.12	3.72	1.8	2.97		-
ADRG11000111	21.55	13.28	21.8	18.36	12.88	13.95	15.62	8.5	14.45		-
ADRG11000027	2.4	1.68	5.6	4.02	2.48	2.04	3.18	0.81	3.55		-
ADRG11000058	84.24	28.04	78.65	95.27	88.98	72.58	39.84	22.31	39.98		-
ADRG11000089	7.81	8.31	10.67	3.58	5.86	4.59	2.92	1.14	2.62	*	-
ADRG11000077	4.08	5.99	6.34	5.76	3.87	4.68	4.23	2	3.08		-
ADRG11000092	41.45	38.67	55.77	73.08	40.91	64.59	24.67	18.9	27.36	*	-
ADRG11000099	43.57	33.01	48.6	58.04	28.24	45.47	24.87	18.08	20.27	*	-
ADRG11000135	52.22	43.8	70.27	58.84	40.95	44.91	35.09	21.07	29.78	*	-
ADRG11000147	6.68	6.78	11.64	6.52	5.38	7	5.38	2.59	5.89		-
ADRG11000159	7.28	7.31	7.55	10.23	10.38	9.89	7.3	4.68	8.62	**	+
ADRG11000160	11.62	12.89	8.28	9.39	13.77	23.6	7.39	5.18	9.59		-
ADRG11000171	4.47	4.72	5.01	4.83	8.43	4.19	3.69	1.77	4.29		-
ADRG11000181	9.73	12.5	18.02	18.65	11.48	13.63	9.96	11.18	7.77		-
BGG111000016	26.13	21.03	22.91	10.68	12.73	20.08	10.32	15.5	18.5	*	-
BGG111000016	63.98	55.25	74.82	18.05	38.94	8.28	17.67	40.85	37.24	*	-
BGG111000017	8.13	8.51	11.48	3.65	5.27	5.54	2.72	4.89	3.8	*	-
BGG111000022	12.88	14.45	15.11	7.84	9.62	8.77	7.97	8.92	10.08	**	-
BGG111000031	13.67	9.32	14.45	9.12	13.53	12.68	6.75	10.28	15.5	*	-
BGG111000042	78.81	92.28	77.71	51.19	53.47	19.5	23.41	16.53	22.63	*	-
BGG111000046	59.14	59.17	43.06	30.88	42.51	35.11	16	15.29	7.23	**	-
BNGH41000025	34.23	58.36	71.87	11.98	22.6	19.45	12.51	15.58	19.08	*	-
BNGH41000026	8.81	10.35	11.92	5.79	4.65	4.41	2.7	3.65	4.11	**	-
BNGH41000027	38.04	28.15	67.91	21.21	34.28	46.33	28.88	28.53	19.37		-
BNGH41000035	71.93	95.4	103.3	77.48	81.38	82.2	85.4	91.4	89.21		-
BNGH41000037	11.37	8.43	16.43	4.88	12.04	10.79	4.26	3.88	5.78		-
BNGH41000042	158.34	222.69	94.88	128.41	120.85	102.56	66.52	26.75	52.39	*	-
BNGH41000046	115.13	80.94	131.08	158.03	89.22	95.02	70.52	58.57	82.66	*	-
BNGH41000056	6.81	7.33	32.48	17.81	11.48	11.22	6.84	4.36	6.34		-
BNGH41000087	7.57	4.78	6.77	6.28	5.74	6.12	2.23	3.11	5.45		-
BNGH41000091	5.88	4.68	7.21	3.92	3.81	2.28	2.51	1.86	5.23	*	-
BNGH41000157	24.78	9.57	39.68	15.95	28.36	14.05	14.96	12.43	25.4		-
BNGH41000159	4.77	2.7	8.76	3.03	4.07	2.47	2.48	1.85	2.4		-
BNGH41000181	15.03	14.18	16.82	9.14	8.43	9.02	9.57	5.48	4.7	**	-
BNGH41000198	5.23	7.17	13.44	2.81	5.92	4	2.64	2.63	2.78		-
BNGH41000219	55.38	63.98	42.98	30.63	34.88	27.18	11.04	4.75	5.11	*	-
BNGH41000229	41.48	41.07	32.45	12.88	20.88	15.08	12.11	29.5	42.2	**	-
BNGH41000237	30.67	28.92	29.88	23.95	17.14	16.35	7.62	11.39	18.2	*	-
BNGH41000238	12.97	8.92	11.13	3.1	5.98	5.05	3.21	2.99	4.83	*	-
BNGH41000243	37.29	22.23	38.12	21.69	20.78	22.28	17.48	13.04	12	*	-
BNGH41000270	7.24	2.74	13.68	5.03	4.81	3.91	2.55	1.86	2.15		-
BRAH11000004	28.95	12.1	22.38	14.77	13.7	15.68	10.21	9.28	9.3		-
BRAH11000016	16.02	18.04	23.1	10.25	11.87	13.62	8.4	6.81	8.28	*	-
BRAH11000021	13.77	12.07	18.61	14.58	14.48	10.85	13.45	8.59	11.48		-
BRAH11000027	4.8	4.82	5.71	4.63	4.59	5.52	2.19	2.04	7.42		-
BRAH11000029	7.2	4.85	5.05	4.87	2.83	2.65	2.93	2.18	1.97	*	-
BRAH11000040	20.85	33.58	27.15	18.04	13.44	12.14	8.98	5.89	6.77	*	-
BRAH11000050	88.78	63.28	107.91	121.47	88.32	92.39	77.36	45.55	64.9		-
BRAH11000051	3.26	3.15	8.62	2.34	1.27	1.81	2.29	1.14	1.62		-
BRAH11000060	103.56	87.24	102.14	122.05	89.97	107.12	58.34	59.07	66.98	**	-
BRAH11000075	6.97	6.63	15	3.29	4.01	3.69	3.52	2.95	3.34		-
BRAH11000081	23.41	12.71	28.79	20.72	20.86	22.65	19.13	9.85	16.04		-
BRAH11000084	219.84	140.27	3.27	8.21	5.55	1.75	108.89	121.2	155.45		-
BRAH11000095	6.87	5.04	6.84	7.47	5.22	4.62	3.81	0.91	1.79	*	-

1/157

【0825】

【表354】

2441

2442

BRAH1000096	5.38	6.14	9.85	5.23	5.99	6.41	5.29	2.17	4.29				
BRAH1000097	32.73	38.25	38.91	16.1	25.45	25.48	34.97	22.26	21.98	*	-		
BRAH1000100	3.44	3.77	6.38	1.33	1.79	1.51	2.06	0.88	1.83	*	*	-	
BRAH1000101	151.17	168.03	162.43	124.31	137.07	112.48	156.93	114.91	72.6	*	-		
BRAH1000104	3.11	5.18	13	2.27	6.18	12.04	5.42	9.52	5.18				
BRAH1000107	4.13	9.11	10.29	5.8	8.98	4.48	6.91	1.75	1.45				
BRAH1000110	63.35	48.25	85.05	53.28	34.71	42.03	42.58	51.83	49.08				
BRAH1000111	17.89	14.24	19.42	8.38	13.57	14.92	11.88	11.38	7.34				
BRAH1000115	7.97	7.93	10.18	8.35	4.56	3.75	5.18	4.92	2.5	*	*	-	
BRAH1000180	13.15	11.62	18.33	11.13	9.11	7.4	12.16	9.59	8.37				
HENBA1000005	35.95	34.82	38.35	20.64	28.11	18.05	30.81	17.47	24.15	**	*	-	
HENBA1000006	9.78	8.8	11.38	1.4	5.53	2.48	3.8	1.71	7.12	**	*	-	
HENBA1000012	142.4	148.44	163.52	138.28	167.18	112.45	92.75	76.02	85.25	**	*	-	
HENBA1000020	190.7	248.05	252	200.63	259.1	189.87	186.01	114.55	161.8				
HENBA1000030	20.38	15.29	27.68	14.24	16.3	11.52	9.94	9.08	14.21				
HENBA1000034	15.22	17.47	15.15	7.38	10.81	14.4	7.69	4.87	8.91	**	-		
HENBA1000042	22.23	17.73	35.18	17.22	20.39	12.68	19.68	13.46	21.05				
HENBA1000045	12.71	14.02	2.48	0.7	7.88	1.89	4.27	2.85	2.17				
HENBA1000046	10.19	13.7	20.59	6.92	10.64	7.67	13.72	12.82	11.87				
HENBA1000047	4.05	7.12	13.48	2.54	4.15	2.97	2.37	0.81	3.55				
HENBA1000048	10.63	14.86	15.61	2.18	5.84	3.23	4.43	0.94	3.67	**	**	-	
HENBA1000050	3.15	3.21	10.5	2.13	3.59	2.35	2.38	0.05	2.24				
HENBA1000051	5.98	3.98	9.78	8.93	5.91	8.48	2.35	1.85	5.68				
HENBA1000058	10.62	11.2	18.18	10.13	8.71	11.8	7.34	2.99	9.18				
HENBA1000072	530.32	492.25	787.01	388.98	521.7	354.83	288.04	398.85	194.47	*	-		
HENBA1000073	26.51	17.12	39.76	19.17	28.84	19.97	20.8	18.82	12.3				
HENBA1000076	30.89	23.42	46.09	20.74	16.14	16.31	22.31	20.51	20.54				
HENBA1000084	48.03	48.38	54.47	61.64	67.48	73.34	43.23	26.25	43.33	*	+		
HENBA1000087	2.35	3.13	9.77	1.66	2.08	0.88	1.69	0.8	1.37				
HENBA1000088	3.05	3.73	12.72	2.97	3.87	2.68	4.68	0.94	2.68				
HENBA1000091	38.82	23.53	75.02	58.3	74.93	71.88	24.1	16.38	30.08				
HENBA1000111	21.39	20.11	44.78	24.89	20.7	28.97	20.62	12.83	18.49				
HENBA1000121	18.15	14.08	45.72	28.88	18.11	19.11	18.09	9.88	11.04				
HENBA1000128	6.74	4.73	14.38	8.16	5.93	6.25	3.31	1.93	2.78				
HENBA1000129	6.94	5.88	13.44	7.51	6.89	6.55	4.13	5.51	2.95				
HENBA1000141	12.4	7.44	22.8	6.83	6.54	5.23	5.19	3.51	3.65				
HENBA1000146	5.55	7.84	15.71	4.82	7.82	7.46	5.3	2.57	3.09				
HENBA1000190	28.41	41.35	48.05	22.83	18.47	23.82	28.68	15.89	24.53	*	-		
HENBA1000154	717.61	698.56	881.96	454.74	531.19	670.47	527.79	542.09	612.33	*	-		
HENBA1000158	10.8	18.5	21	9.98	4.23	6.43	3.69	3.2	3.38	*	-		
HENBA1000158	18.74	13.45	14.16	7.7	2.81	4.64	5.39	2.06	3.28	**	**	-	
HENBA1000188	8.01	6.75	7.08	5.88	4.99	5.88	6.88	2.42	4.44	*	-		
HENBA1000180	3.02	3.28	4.77	4.2	2.93	2.73	2.69	1.65	3.39				
HENBA1000185	19.4	18.45	23.05	27.72	15.45	18.55	13.51	10.97	19.3				
HENBA1000188	8.28	5.61	7.48	5.82	8.91	3.78	3.34	2.85	4.68	*	-		
HENBA1000193	7.68	7.79	11.97	13.74	6.68	7.85	5.99	5.1	6.65				
HENBA1000194	31.21	25.82	29.1	20.52	12.15	13.24	9	11.16	12.72	*	**	-	
HENBA1000201	19.81	8.63	14.31	11.15	8.36	7.64	5.43	4.05	4.24	*	-		
HENBA1000213	2.97	4.59	8.02	5.38	2.78	4.45	3.2	1.24	3.2				
HENBA1000216	7.82	8.45	11.73	8.7	8.98	8.82	8.13	4.35	5.58	*	-		
HENBA1000227	2.78	4.56	8.5	4.18	3.58	2.84	3.35	1.71	2.27				
HENBA1000231	24.52	27.31	24.37	15.72	17.43	21.18	21.29	12.6	20.48	*	-		
HENBA1000237	78.81	74.32	83.86	68.84	89.97	42.93	44.61	22.45	38.37	**	-		
HENBA1000243	13.82	10.47	25.88	21.77	11.3	17.41	12.09	8.84	11.38				
HENBA1000244	9	7.04	9.25	7.59	2.88	4.81	3.47	4.91	3.18	**	-		
HENBA1000251	4.62	4.41	5.87	4.22	1.83	3.25	2.37	2.22	2.32	**	-		
HENBA1000254	10.09	13.73	13.57	7.22	5.82	7.35	8.91	6.98	9.5	*	*	-	
HENBA1000254	5.1	3.58	11.18	4.22	2.15	2.47	2.98	1.73	5.35				
HENBA1000259	4.34	6.24	5.83	3.94	4.73	5.35	3.13	1.23	3.18	*	-		
HENBA1000275	22.4	40.99	22.83	21.15	22.57	23.38	10.3	5.45	12.41	*	-		
HENBA1000280	6.51	8.25	15.12	7.28	7.37	8.84	7.59	2.9	7.78				
HENBA1000282	18.71	16.98	35.85	19.88	12.71	13.87	18.74	14.02	21.9				
HENBA1000287	10.5	12.84	12.5	3.76	7.11	8.61	3.4	8.14	8.93	**	-		
HENBA1000288	14.15	21.12	38.73	13.92	11.1	18.3	14.48	11.79	19.51				

2 / 157

2443

2444

HENBA1000290	5.54	5.24	10.08	5.32	1.73	3.05	3.54	2.47	2.22				
HENBA1000296	9.5	12.84	20.78	5.15	12.54	6.88	5.85	5.2	3.4	*		-	
HENBA1000300	35.72	34.39	45.63	27	31.39	25.7	28.91	18.61	28.82	*		-	
HENBA1000302	3.82	6.7	8.9	1.71	5.14	2.38	3.51	1.14	3.24				
HENBA1000303	7.88	7.6	13.78	3.6	4.76	4.02	4.54	2.89	3.82	*		-	
HENBA1000304	20.77	16.02	19.73	12.6	11.48	11.85	12.2	13.31	13.14	**	*	-	
HENBA1000307	7.67	8.03	6.72	5.49	4.18	4.06	1.61	1.61	3.87	**	**	-	
HENBA1000312	26.83	27.01	26.09	21.32	17.64	18.68	18.71	13.82	23.64	*		-	
HENBA1000318	14.81	10.63	16.92	6.25	7.77	6.66	2.82	1.83	3.66	*	**	-	
HENBA1000327	16.1	18.58	31.73	13.19	19.68	35.1	18.48	11.24	13.77				
HENBA1000333	4.8	4.82	8.3	2.28	3.1	2.42	2.41	0.71	1.67	*	*	-	
HENBA1000338	14.85	18.8	25.1	10	13.67	12.74	13.29	10.79	10.59				
HENBA1000343	8.78	12.6	14.72	7.21	10.87	6.74	6.68	5.91	8.78	*		-	
HENBA1000349	7.48	8.48	16.3	4.37	4.58	5.11	3.79	2.9	11.15				
HENBA1000351	16.95	12.69	26.79	18.1	17.65	14.44	11.22	7.92	17.2				
HENBA1000355	11.89	7.38	8.03	7.48	3.22	2.44	1.27	2.39	3.56	*		-	
HENBA1000356	34.47	35.59	32.24	39.82	29.15	32.82	22.27	24.67	20.29	**		-	
HENBA1000357	16.16	16.75	30.37	11.91	12.99	11.48	14.62	9.28	12.57				
HENBA1000368	10.73	7.46	15.42	5.17	5.07	3.55	5.42	3.22	6.53	*		-	
HENBA1000369	10.72	11.69	16.34	4.55	5.5	4.39	6.78	3.13	5.67	*	*	-	
HENBA1000370	9	13.55	13.78	4.72	8.24	7.37	5.92	3.13	3.05	*	*	-	
HENBA1000378	90.89	73.26	0.51	1.99	5.8	2.22	56.72	52.09	77.09				
HENBA1000387	25.37	30.02	2.08	0.67	28.79	0	22.74	18.21	28.75				
HENBA1000388	16.44	15.64	23.44	8.78	10.05	16.24	16.41	8.42	9.98				
HENBA1000390	10	6.3	9.6	4.59	5.15	2.55	7.47	3.11	8.98	*		-	
HENBA1000392	7.7	8.4	14.19	5.19	4.42	2.87	4.91	3.87	4.75				
HENBA1000396	9.8	6.67	9.18	1.73	1.25	0.34	3.88	1.99	4.11	**	*	-	
HENBA1000411	4.32	4.84	13.72	2.13	3.17	1.91	3.38	1.45	2.17				
HENBA1000418	18.48	14.65	17.48	5.45	8.2	5.71	4.58	2.35	3.71	*	**	-	
HENBA1000422	9.88	6.28	0	0.69	0.58	0.28	3.82	3.45	7.58				
HENBA1000428	6.49	7.69	1.25	0.25	6.33	0	5.22	5.3	9.15				
HENBA1000434	6.78	3.73	4.63	2.87	2.1	1.84	5.95	1.48	3.75	*		-	
HENBA1000442	3.73	3.34	6.65	2.11	2.23	1.05	2.87	2.62	6.58				
HENBA1000443	6.03	7.17	14.31	3.08	6.26	3.05	8.49	4.27	7.22				
HENBA1000448	58.88	47.2	7.47	1.38	1.35	0.83	46.56	24.3	38.67	*		-	
HENBA1000456	14.88	20.85	28.05	6.95	15.88	10.1	10.12	2.72	7.69	*		-	
HENBA1000459	15.34	18.32	27.74	11	17.98	14.9	11.74	7.37	11.08				
HENBA1000460	23.55	18.11	28.93	19.24	20.63	19.37	10.12	7.72	17.35				
HENBA1000462	25.82	7.48	2.25	1.21	3.05	10.25	13.12	8.17	7.37				
HENBA1000484	7.58	4.61	6.98	3.88	3.64	2.38	3.55	1.28	2.63	*	*	-	
HENBA1000488	3	4	6.33	1.85	2.29	0.39	1.45	0.17	1.64	*		-	
HENBA1000489	22.54	28.03	34.83	17.42	17.27	14.02	27.85	14.61	18.23	*		-	
HENBA1000477	17.38	15.06	7.59	1.07	12.5	0.35	11.11	7.11	9.18				
HENBA1000481	63.47	60.18	49.05	5.51	28	7.01	17.9	5.67	5.55	*	**	-	
HENBA1000488	14.11	13.35	8.89	0.4	9.3	4.88	6.34	1.81	3.64	*	*	-	
HENBA1000480	9	6.13	11.44	3.85	4.12	5.04	6.84	2.63	7.56	*		-	
HENBA1000491	5.32	6.61	17.42	8.34	8.08	9.74	8.93	4.28	9.22				
HENBA1000488	14.5	18.25	24.08	10.95	15.11	14.4	16.82	10.93	14.38				
HENBA1000501	7.87	9.11	2.74	0.21	2.74	4.51	8.04	4.62	6.97				
HENBA1000504	12.64	5.58	11.75	3.41	4.13	2.59	2.67	4.18	4.58	*		-	
HENBA1000505	6.92	8.64	7.68	0	1.28	6.84	6.6	2.13	4.7	**	*	-	
HENBA1000507	0.45	1.66	6.28	0	1.2	0.18	0.77	0	0.38				
HENBA1000508	28.91	34.97	60.59	22.03	32.11	27.29	38.18	21.05	20.98				
HENBA1000518	3.76	3.28	6.48	5.32	4.44	6.01	2.59	1.88	3.38				
HENBA1000519	85.07	81.88	141.09	42.18	50.11	71.14	47.48	35.9	57.59	*		-	
HENBA1000520	6.49	7.75	13.66	6.32	6.26	4.22	5.77	4.5	4.14				
HENBA1000523	11.85	6.14	1.44	0	1.42	0.29	4.15	3.68	2.35				
HENBA1000531	8.26	6.64	16	4.73	9.37	11.48	9.54	4.57	6.14				
HENBA1000534	28.94	37.16	55.85	24.81	22.78	20.48	41.37	28.08	32.84				
HENBA1000538	14.97	19.18	30.43	12.11	14.95	11.04	13.25	9.03	10.27				
HENBA1000540	18.41	10.82	26.42	7.05	8.02	12.43	11.49	6.58	4.88				
HENBA1000542	68.21	65.88	94.9	105.13	128.6	107.62	49.4	38.5	60.6	*	*	+	-
HENBA1000545	10.72	8.72	19.53	13.15	9.91	19.84	7.59	4.75	4.2				
HENBA1000547	16.09	12.61	26.19	11.54	13.04	5.83	5.77	5.14		*		-	

3 / 157

2446

HENBA1000551	81.43	29.13	74.82	42.08	38.34	36.48	27.97	18.18	17.63
HENBA1000555	23.07	51.65	28.69	16.76	19.18	84.47	21.17	21.24	16.53
HENBA1000567	20.20	22.07	26.42	16.84	15.85	18.08	21.8	11.99	15.53
HENBA1000581	0.79	6.3	13.7	3.18	1.89	4.15	4.27	2.09	3.68
HENBA1000583	2.71	6.02	9.85	3.7	6.59	3.42	4.89	1.88	4.27
HENBA1000587	2.91	1.65	3.34	2.35	0.99	3.67	2.88	1.41	2.17
HENBA1000588	8.19	5.62	14.84	6.57	7.03	7.81	7.88	3.61	9.31
HENBA1000589	2.57	3.31	7.07	4.84	2.77	3.17	2.3	1.18	2.24
HENBA1000595	24.44	15.32	39.5	24.17	18.92	19.53	17.36	14	17.06
HENBA1000598	7.1	5.81	8.91	7.19	8.88	8.85	8.33	2.25	5.9
HENBA1000600	5.23	6.08	5.7	5.38	6.93	6.68	4.07	2.31	4.39
HENBA1000611	24.12	22.46	26.56	10.74	18.87	21.61	11.03	6.78	13.01
HENBA1000612	30.95	41.43	24.63	10.67	31.41	23.82	10.57	11.36	12.87
HENBA1000614	5.11	7.29	6.67	8.37	4.71	6.88	5.32	3.27	3.99
HENBA1000624	21.22	17.48	22.7	28.09	18.45	20.89	19.32	9.43	17.26
HENBA1000637	59.03	59	51.96	71.67	51.82	56.93	60.28	42.6	51.42
HENBA1000648	9.41	4.42	6.41	8.42	6.12	4.87	9.82	3.99	8.94
HENBA1000672	7.92	7.76	16.15	9.11	9.27	13.99	7.94	3.81	11.14
HENBA1000934	52.78	74.71	70.78	38.68	65.82	64.7	32.15	24.17	25.34
HENBA1000936	22.12	28.11	26.27	26.4	22.98	20.49	9.12	8.29	12.36
HENBA1000937	8.88	11.75	7.87	5.21	9.28	6.88	4.93	3.24	3.68
HENBA1000955	30.73	31.89	54.9	27.54	18.27	22.75	21.5	16.39	24.41
HENBA1000957	14.08	9.28	10.28	5.98	7.53	6.3	3.93	18.82	5.24
HENBA1000962	6.27	5.79	10.27	3.02	6.27	5.29	2.93	4.16	3.38
HENBA1000964	6.20	6.77	9.71	6.57	5.26	8.62	2.22	2.8	4.62
HENBA1000971	30.41	19.43	27.66	31.56	22.01	24.63	20.58	14.98	15.58
HENBA1000973	24.11	24.74	28.83	18.72	18.93	18.52	21.96	17.14	28.31
HENBA1000975	12.22	13.93	16.24	11.01	14.94	12.36	11.46	6.91	7.37
HENBA1000978	5.24	8.15	11.3	5.35	7.05	6.17	5.02	1.55	1.87
HENBA1000982	64.67	35.39	63.17	14.17	18.8	19.86	13.73	23.78	33.18
HENBA1000986	21.71	40.16	25.85	10.91	16.11	14.54	4.3	11.52	8.5
HENBA1000992	10.84	17.35	25.78	7.39	11.26	13.13	7.65	7.08	9.42
HENBA1000995	4.8	3.75	8.97	3.21	1.09	3.1	3.08	4.72	3.51
HENBA1000999	10.43	11	19.66	6.79	9.41	8.35	11.72	8.05	13.37
HENBA1000999	27.85	31.72	34.47	18.34	17.98	21.82	22.89	15.55	20.61
HENBA1000999	21.36	28.6	18.8	18.63	23.29	21.74	21.77	16.83	13.69
HENBA1000999	12.12	14.07	15.25	12.43	11.9	8.78	7.24	4.41	3.48
HENBA1000999	32.16	32.06	69.89	34.36	26.18	30.36	38.32	22.04	41.04
HENBA1000999	27.97	25.82	19.68	6.88	11.41	11.31	4.45	8.11	7.09
HENBA1000999	8.15	9.73	16.83	6.74	7.08	9.59	5.57	4.8	5.38
HENBA1000999	9.38	7.92	11.45	7.38	12.77	8.49	9.33	8.94	12.57
HENBA1000999	7.76	5.78	11.72	5.12	7.77	27.92	5.16	3.39	8.47
HENBA1000999	0.81	6.54	13.88	4.04	5.62	4.48	4.88	2.87	4.45
HENBA1000999	5.32	4.72	12.62	2.69	5.68	3.35	3.85	2.1	3.4
HENBA1000999	7.08	8.97	13.44	6.2	9.91	9.41	8.29	3.96	9.55
HENBA1000999	18.03	20.17	33.28	13.55	12.62	15.47	12.49	12.59	18.78
HENBA1000999	9.61	11.01	10.43	7	5.88	6.12	4.8	4.54	5.88
HENBA1000999	14.73	8.44	14.98	4.78	5.04	11.61	6.39	6.15	8.17
HENBA1000999	11.82	8.81	16.04	4.91	5.14	4.63	2.58	3.42	2.84
HENBA1000999	13.61	13.09	20.89	7.09	10.48	10.58	6.68	4.5	6.77
HENBA1000999	7.76	13.89	13.94	4.47	6.28	7.04	9.68	6.14	7.88
HENBA1000999	3.65	4.13	15.13	0.99	5.68	8.11	2.71	1.47	3.79
HENBA1000999	21.34	27.53	42.71	22.13	28.3	22.38	30.47	15.88	17.68
HENBA1000999	8.15	5.35	9.22	6.03	4.9	5.38	2.5	2.74	2.78
HENBA1000999	8.64	7.55	8.23	4.94	6.39	5.68	3.78	3.65	3.48
HENBA1000999	29.92	28.02	44.77	28.48	22.57	21.5	32.85	19.21	32.35
HENBA1000999	30.39	19.19	19.38	8.84	18.27	15.1	13.32	16.07	12.64
HENBA1000999	4.74	2.72	8.7	3.43	2.84	2.21	2.09	1.11	2.74
HENBA1000999	46.02	47.69	50.56	27.4	35.01	30.28	30.89	15.81	24.68
HENBA1000999	12.65	12.98	18.93	6.72	11.22	3.95	2.79	1.89	3.94
HENBA1000999	6.54	10.54	13.74	4.86	7.54	6.86	4.29	1.48	3.18
HENBA1000999	13.88	14.84	28.11	13.31	20.08	16.37	7.68	5.83	12.73
HENBA1000999	12.7	15.8	16.64	6.55	5.66	4.68	3.19	1.78	1.97
HENBA1000999	116.36	117.03	138.17	175.69	117.43	151.33	112.02	74.79	83.7

2447

2448

HENBA1000843	12.55	15.69	27.26	12.49	12.38	9.92	12.87	6.01	13.01				
HENBA1000851	7.99	8.88	11.08	3.31	3.75	3.94	6.33	2.77	2.83	**	*	-	-
HENBA1000852	16.57	13.81	23	7.9	10.22	6.68	11.13	8.47	9.44	*	*	-	-
HENBA1000857	8.41	40.09	21.59	7.82	12.17	12.73	16.16	8.68	5.82				
HENBA1000859	6.66	5.92	14.84	4.94	8.68	4.87	5.86	1.59	2.82				
HENBA1000870	16.63	14.8	15.74	8.74	7.25	8.9	6.35	3.09	4.06	**	**	-	-
HENBA1000872	19	16.2	19.95	12.25	11.2	8.26	7.42	4.77	5.4	**	**	-	-
HENBA1000875	7.38	6.41	11.16	2.07	4.19	4.82	3.98	2.9	2.7				
HENBA1000876	12.29	12.13	19.01	9.58	10.69	6.67	12.67	8.69	9.73				
HENBA1000907	4.95	12.04	11.83	5.83	7.68	10.61	8.24	2.82	6.62				
HENBA1000908	7.89	3.54	11.79	2.7	10.88	1.89	2.83	0.62	3.13				
HENBA1000910	4.91	8.05	12.14	3.81	5.91	4.66	5.84	3.7	4.74				
HENBA1000918	21.04	31.68	45	25.17	23.52	19.82	25.88	9.86	18.68				
HENBA1000919	4.44	2.5	5.8	3.35	2.88	3.17	3.08	0.43	2.37				
HENBA1000934	4.04	4.57	5.7	3.99	3.21	2.45	2.55	1.64	2.64	*		-	-
HENBA1000935	4.4	5.61	6.39	3.11	4.92	2.1	3.77	2.49	2.75	*		-	-
HENBA1000940	8.69	8.39	12.11	8.45	5.94	4.28	6.33	4.58	4.22	*	*	-	-
HENBA1000942	12.05	15.91	20.05	9.92	11.92	8.31	14.27	9.3	8.99				
HENBA1000943	2.81	3.51	9.04	0.06	1.83	0.43	1.65	0.02	0.85				
HENBA1000946	5.48	9.79	16.74	3.82	4.51	6.06	3.38	0.73	4.11				
HENBA1000960	20.15	53.34	82.32	44.07	41.68	37.38	55.19	30.38	32.98				
HENBA1000962	10.91	13.68	14.89	4.87	9.87	8.15	5	5.38	7.85	*	**	-	-
HENBA1000968	9.18	5.88	12.63	7.34	6.67	5.5	4.92	3.18	5.66				
HENBA1000971	12.8	9.28	22.24	4.73	7.24	3.99	4.58	3.74	2.56	*		-	-
HENBA1000972	8.17	5.02	15.82	6.78	5.68	6.86	5.46	4.31	5.77				
HENBA1000974	17.98	18.06	33.14	22.9	19.88	10.01	13.84	8.08	13.41				
HENBA1000975	8.54	8.18	15.36	5.78	4.48	4.06	6.5	5	4.65				
HENBA1000979	12.05	14.89	32.47	11.88	10.02	10.58	16.5	7.78	11.37				
HENBA1000981	31.09	24.35	80.14	12.11	18.86	23.49	28.22	14.59	14.47				
HENBA1000983	18.01	16.07	24.38	8.83	12.27	20.3	15.26	9.78	18.85				
HENBA1000985	7.28	3.65	6.88	4.91	4.64	4.99	4.05	3.11	1.91				
HENBA1000986	13.51	10.48	26.88	15.8	12.6	15.39	12.22	6.19	7.06				
HENBA1000991	15.73	13.88	24.7	13.38	13.44	8.85	12.82	7.37	7.8				
HENBA1001007	6.2	4.4	10.1	3.64	3.86	2.18	2.52	1.45	2.54				
HENBA1001008	5.26	4.16	10.22	2.95	3.18	4.11	4.27	1.48	2.86				
HENBA1001009	2.68	2.47	7.99	2.44	3.63	2.07	2.06	0.9	2.84				
HENBA1001014	36.78	43.69	57.32	33.05	32.39	34.29	37.95	16.24	32.58				
HENBA1001017	7.14	5.71	12.9	9.32	6.72	6.57	4.21	2.97	4.79				
HENBA1001019	6.38	3.7	5.48	7.22	3.14	5.53	3.87	2.27	2.61				
HENBA1001020	9.13	7.41	10.99	8.19	4.18	5.07	8.01	5.07	4.27				
HENBA1001021	4.2	4.32	9.05	3.87	3.59	5.88	6.3	1.95	6.15				
HENBA1001022	10.5	4.41	10.28	9.91	9.22	14.59	10.15	4.7	6.8				
HENBA1001024	1.82	2.83	5.27	3.68	2.85	3.77	2.92	1.3	2.29				
HENBA1001026	3.74	3.71	5.2	2.99	4.11	4.69	3.01	1.9	3.66				
HENBA1001043	6.37	5.28	6.61	4.42	5.68	7.08	4.73	0.65	3.01				
HENBA1001051	120.57	138.21	211.05	111.88	82.79	89.92	82.72	88.42	94.85				
HENBA1001052	6.89	6.44	5.5	6.15	5.41	3.95	3.07	3.65	4.06	**		-	-
HENBA1001059	56.7	56.44	78.92	98.8	52.77	73	38.68	40.32	41.94	*		-	-
HENBA1001060	22.93	14.44	26.88	18.88	16.28	14.69	14.87	11.27	12.22				
HENBA1001064	7.94	5.9	11.63	5.51	6.84	7.38	5.01	1.85	2.88				
HENBA1001071	2.97	5.22	8.35	8.83	8.39	5.39	3.28	1.85	3.81				
HENBA1001077	6.95	10.47	5.69	10.99	11.81	13.69	3.82	3.2	5.87				
HENBA1001078	13.55	16.29	18.17	15.7	24.22	16.62	14.25	7.88	7.1				
HENBA1001080	18.28	14.99	25.65	14.28	8.2	7.86	14.01	11.12	10.13				
HENBA1001084	10.12	12.02	19.64	11.43	8.23	10.4	8.52	5.91	8.47				
HENBA1001085	12.67	14.13	17.48	10.37	7.16	7.36	9.72	8.19	7.2	*	*	-	-
HENBA1001088	3.47	2.77	6.21	3.04	2.22	2.64	4.71	4.47	2.67				
HENBA1001093	5.89	7.48	12.08	5.52	4.12	4.26	4.75	3.73	5.78				
HENBA1001094	3.87	3.67	7.3	1.83	2.98	1.32	1.79	1.61	3.04				
HENBA1001099	4.27	4.88	10.87	6.66	5.88	4.18	3.44	1.48	3.72				
HENBA1001104	3.23	5.58	9.6	7.86	21.55	10.79	5.83	1.68	4.13				
HENBA1001109	55.09	49.81	96.04	54.28	41.81	45.47	44.91	14.62	57.09				
HENBA1001114	101.03	180.87	118.81	64.42	78.99	89.5	43.16	105.03	85.05				
HENBA1001121	8.11	7.04	9.73	5.44	4.04	4.32	4.43	2.6	4.72	*	*	-	-

5 / 157

2449

2450

HENBA1001122	8.39	9.29	17.71	6.58	5.73	6.53	2.48	3.1	3.28	*	-
HENBA1001123	17.13	17.37	24.47	15.29	11.78	13.71	12.33	9.75	10.61	*	-
HENBA1001133	8.92	6.97	12.83	3.89	4.11	4.57	3.05	2.9	3.3	*	-
HENBA1001137	8.57	7.37	12.68	4.24	4.79	2.86	4.1	2.06	3.89	*	-
HENBA1001140	9.64	11.4	16.66	10.87	12.26	10.82	8.45	8.07	5.83	*	-
HENBA1001144	75.62	79.41	130.35	64.7	57.88	71.91	43.23	35.45	52.04	*	-
HENBA1001145	228.88	419.26	287.93	34.88	83.53	106.81	28.23	79.02	93.39	*	-
HENBA1001158	11.25	10.45	12.32	5.71	7.87	10.11	6.5	4.55	5.45	*	-
HENBA1001172	13.23	15.14	21.25	10.8	9.67	9.91	10.54	8.89	11.23	*	-
HENBA1001174	4.77	4.96	13.83	3.36	4.72	5.71	5.53	3.53	3.57	*	-
HENBA1001175	25.84	49.85	37.18	5.53	38.99	18.9	6.53	8.52	5.43	*	-
HENBA1001182	253.75	199.19	217.61	213.83	179.08	196.11	166.5	128.3	113.47	*	-
HENBA1001184	2.72	5.66	8.48	2.92	7.84	3.84	2.54	1.64	3.55	*	-
HENBA1001192	7.71	9.61	13.89	5.11	4.84	2.98	2.49	2.97	4.44	*	-
HENBA1001196	20.51	33.03	45	22.81	18.19	24.13	7.47	12.08	16.7	*	-
HENBA1001197	377.82	621.74	482.95	613.89	355.35	482.48	223.5	509.73	494.09	*	-
HENBA1001208	6.77	4.91	9.09	4.11	4.25	4.69	3.47	2.44	4.54	*	-
HENBA1001213	162.45	150.13	151.68	224.78	129.48	188.81	135.18	97.08	115.8	*	-
HENBA1001214	10.4	15.18	18.48	6.51	7.45	6.56	4.99	3.77	6.34	*	-
HENBA1001221	12.82	12.25	19.21	2.1	6.95	5.45	2.31	0.75	2.8	*	-
HENBA1001225	4.23	4.43	8.6	1.43	2.73	4.08	2.02	0.94	2.52	*	-
HENBA1001225	43.15	51.42	64.65	33.28	48.87	38.83	29.67	18.32	24.38	*	-
HENBA1001228	5.8	4.52	8.07	5.68	4.51	3.32	3.32	1.37	1.24	*	-
HENBA1001229	171.48	141.71	227.85	215.78	153.09	225.68	105.81	85.9	109.68	*	-
HENBA1001235	51.44	62.4	68.51	46.91	34.3	36.65	26.85	20.23	22.59	*	-
HENBA1001236	19.14	19.72	19.8	10.88	15.88	9.5	15.04	10.36	18.33	*	-
HENBA1001242	153.09	161.43	160.67	152.68	145.07	151.85	99.19	75.78	86.89	*	-
HENBA1001247	33.3	29.15	48.14	21.32	32.34	22.28	24.27	20.52	27.05	*	-
HENBA1001253	159.45	91.8	142.81	118.02	138.28	127.67	95.67	74.41	74.86	*	-
HENBA1001257	10.7	11.57	12.78	5.67	10.33	4.37	9.19	6.06	4.62	*	-
HENBA1001261	6.83	4.18	8.42	7.74	2.14	4.76	3.41	1.78	5.45	*	-
HENBA1001262	12	7.56	5.09	4.62	4.49	6.27	3.32	1.44	4.7	*	-
HENBA1001265	15.44	18.41	25	9.71	8.97	12.14	13.41	14.3	11.72	*	-
HENBA1001268	22.75	28.89	30.68	18.84	17.03	12.35	18.86	10.28	14.84	*	-
HENBA1001269	61.54	134.76	5.56	0.22	1.58	0.07	24.28	23.59	16.64	*	-
HENBA1001272	5.33	4.58	13.33	2.87	3.84	5.83	8.1	1.87	1.85	*	-
HENBA1001279	26.1	31.13	42.25	28.49	33.02	28.73	22.53	17.36	24.84	*	-
HENBA1001281	45.22	55.86	55.89	31.39	34.2	28.87	21.08	11.24	15.62	*	-
HENBA1001284	42.89	33.78	59.69	41.48	38.84	48.02	50.11	34.66	41.6	*	-
HENBA1001289	4.71	4.83	5.28	3.13	4.27	3.23	3.58	1.6	3.12	*	-
HENBA1001291	35.57	24.72	38.55	19.37	18.57	13.13	19.39	15.96	11.69	*	-
HENBA1001294	8.41	15.41	15.09	9.8	8.09	6.28	18.25	5.05	15.73	*	-
HENBA1001296	7.87	20.06	12.89	5.33	6.68	3.52	5.19	2.03	4.83	*	-
HENBA1001297	10.62	19.22	19.18	4.41	7.4	5.61	5.49	1.83	3.17	*	-
HENBA1001299	34.22	39.64	53.21	25.14	29.23	20.68	38.68	22.86	32.79	*	-
HENBA1001302	25.32	40.78	38.96	17.94	26.16	22.11	18.04	11.91	7.95	*	-
HENBA1001303	10.97	8.86	13.52	4.53	5.04	4.42	4.71	4.03	7.45	*	-
HENBA1001306	14.82	11.94	17.26	5.51	7.45	8.38	3.81	5.28	4.41	*	-
HENBA1001308	44.73	35.49	68.19	29.34	49.52	28	40.25	28.31	38	*	-
HENBA1001310	8.14	8.04	8.8	2.67	3.81	3.08	5.37	3.84	2.99	*	-
HENBA1001312	42.22	39.38	44.67	10.33	40.14	18.42	38.81	18.98	13.33	*	-
HENBA1001319	3.83	6.75	12.77	3	5.03	19.5	4.72	0.95	4.62	*	-
HENBA1001322	10.16	10.91	17.38	3.57	7.15	4.09	5.93	2.59	3.38	*	-
HENBA1001323	41.14	38.93	47.79	17.38	18.24	15.31	13.85	10.42	3.9	*	-
HENBA1001326	38.28	32.16	36.43	4.89	17.62	29.01	7.49	3.64	9.8	*	-
HENBA1001327	22.04	9.71	16.11	7.86	12.59	16.77	4.84	4.12	2.76	*	-
HENBA1001330	29.76	31.78	64.11	34.57	26.99	26.81	34.76	20.01	20.9	*	-
HENBA1001348	8.77	6.35	12.87	5.82	7.19	5.23	5.41	4.95	2.78	*	-
HENBA1001350	11.78	12.52	19.81	3.72	3.89	6.69	7.45	5.17	4.88	*	-
HENBA1001351	31.73	33.01	88.64	18.41	14.04	17.67	14.45	12.68	8.51	*	-
HENBA1001352	18.11	10.31	24.58	3.8	5.27	4.13	8.48	3.75	2.44	*	-
HENBA1001353	239.33	87.11	198.31	30.75	66.77	70.17	148.92	56.48	39.11	*	-
HENBA1001358	74.35	56.7	93	21.23	34.67	61.64	20.12	8.88	14.9	*	-
HENBA1001361	9.19	10.7	19.27	11.26	10.32	15.03	5.58	3.47	4.41	*	-

6/157

2451

2452

HEMBA1001364	5.59	5.71	8.25	5.34	8.32	8.23	4.19	2.87	3.41	*	-
HEMBA1001375	10.82	8.04	9.98	6.58	6.52	8.55	3.93	8.03	4.02		
HEMBA1001377	30.09	30.45	47.44	34.01	31.38	34.79	26.34	19.08	22.43		
HEMBA1001383	3.91	6.12	10.35	4.08	4.49	3.29	3.04	1.16	2.13		
HEMBA1001387	4.6	4.17	9.82	4.35	3.61	5.27	5	2.3	4.29		
HEMBA1001388	3.38	12.99	10.77	8.84	12.96	11.27	14.15	2.79	4.79		
HEMBA1001390	18.62	20.96	41.58	20.9	20.32	29.35	22.75	17.7	27.58		
HEMBA1001391	2.64	5.63	8.65	8.35	4.1	4.57	9.4	1.58	2.14		
HEMBA1001398	7.95	9	20.79	14.55	10.11	10.1	9.09	3.85	9.98		
HEMBA1001405	12.88	15.37	29.38	17.27	15.08	18.39	13.42	7.68	15.76		
HEMBA1001406	1.07	1.48	8.07	4.84	2.23	2.45	3.84	0.63	2.8		
HEMBA1001407	26.07	42.48	69.39	34.7	28.94	28.07	26.15	15.14	21.98		
HEMBA1001411	12.19	10.95	23.76	12.34	12.81	18.6	13.01	4.8	11.91		
HEMBA1001413	3.88	7.52	13.68	7.68	4.74	6.7	4.77	2.46	3.91		
HEMBA1001414	10.32	10.99	22.49	13.09	11.95	9.01	7.42	9.57	11.87		
HEMBA1001415	108.07	112.24	218.31	263.58	139.52	200.62	94.4	74.84	97.27		
HEMBA1001416	5.94	4.6	7.69	5.08	3.57	4.15	3.74	1.73	3.21	*	-
HEMBA1001432	3.3	3.49	7.34	4.93	5.95	7.57	4.22	3.23	2.6		
HEMBA1001433	5.11	5.55	7.48	5.92	7.1	8.22	4.04	2.29	5.52		
HEMBA1001435	443.81	559.24	661.15	777.73	679.62	850.28	840.12	377.73	869.08		
HEMBA1001442	6.84	10.08	7.28	7.83	12.04	8.48	5.85	1.98	2.4		
HEMBA1001448	3.77	5.34	8.49	8.2	4.64	4.7	3.45	2.22	3.65		
HEMBA1001450	87.92	103.44	160.6	292.41	124.8	175.34	89.47	74.72	92.31		
HEMBA1001454	32.58	33.13	59.47	19.79	12.7	25.66	18.88	24.9	33.22		
HEMBA1001455	4.27	8.3	7.42	3.13	6.27	4.98	6.18	1.51	8.98		
HEMBA1001459	13.65	28.28	34.13	12.19	8.69	14.19	9.18	11.22	13.6		
HEMBA1001461	30.98	26.88	43.29	83.45	32.08	38.73	26.62	18.12	23.29		
HEMBA1001482	8.43	9.05	11.64	4.2	7.04	5.83	4.19	2.77	3.22	*	**
HEMBA1001483	1.15	4.7	4.74	4.99	6.6	4.02	4.68	1.57	3.48		
HEMBA1001469	23.94	42.87	64.78	64.17	33.38	52.34	14.36	18.57	20.01		
HEMBA1001473	7.37	9.89	11.35	4.59	3.3	4.44	2.71	5.14	2.51	*	*
HEMBA1001477	143.72	139.96	283.29	91.65	145.02	139.05	125.85	114.31	83.92		
HEMBA1001478	3.38	2.99	11.83	2.83	2.55	3.35	2.27	2.19	2.08		
HEMBA1001480	12.29	13.63	23.93	4.88	9.2	12.63	3.97	2.72	4.12	*	-
HEMBA1001483	20.97	27.37	23.32	22.22	18.94	28.05	14.39	10.01	18.51	*	-
HEMBA1001490	14.01	20.37	28.13	15.25	17.46	15.48	22.44	13.02	12.8		
HEMBA1001495	29.67	31.18	55.68	75.52	52.85	89.44	21.73	15.8	24.83		
HEMBA1001497	60.34	63.13	114.17	128.19	117.89	137.3	43.07	37.4	51.44		
HEMBA1001510	6.77	7.43	13.73	8.69	8.1	7.82	5.34	3.9	10.87		
HEMBA1001515	4.98	2.98	9.2	1.95	4	3.02	1.96	0.89	7.49		
HEMBA1001517	8.97	7.98	15.08	6.1	6.94	5.43	7.19	5.25	8.47		
HEMBA1001522	12.78	16.29	21.22	10.39	13.01	11	10.54	9.25	10.68		
HEMBA1001525	8.05	10.88	20.31	6.19	10.35	7.24	6.93	4.75	7.32		
HEMBA1001533	10.87	13.52	23.12	12.09	15.84	9.55	9.42	6.16	11.58		
HEMBA1001547	5.91	8.31	10.54	8.17	5.66	9.57	6.23	2.5	7.38		
HEMBA1001552	2.97	4.72	5.75	6.36	5.58	2.49	2.58	1	6.09		
HEMBA1001553	11.17	10.19	22.05	10.88	12.27	11.05	5.91	7.82	10.68		
HEMBA1001557	7.49	2.45	11.09	3.83	4.84	6.27	3.14	6.97	8.01		
HEMBA1001563	8.8	2.01	15.37	5.55	6.29	6.53	4.15	2	8.58		
HEMBA1001566	4.27	3.01	8.62	2.19	3.8	3.87	2.2	1.62	3.68		
HEMBA1001569	9.15	15.85	27.87	11.41	11.78	14.08	8.02	5.27	11.03		
HEMBA1001570	10.89	22.06	19.57	12.99	13.03	18.53	9.14	4.13	3.81	*	-
HEMBA1001579	85.92	121.25	222.35	134.48	102.08	79.98	25.8	43.38	73.16		
HEMBA1001581	4.11	4.25	7.71	3.25	2.45	2.33	0.91	1.64	3.55		
HEMBA1001582	12.19	16.67	20.04	15.87	10.82	11.89	13.88	8.89	21.33		
HEMBA1001585	8.81	4.99	13.1	7.78	5	7.18	4.04	1.85	5.57		
HEMBA1001589	7.95	8.96	19.02	7.88	6.17	6	7.81	5.42	8.45		
HEMBA1001595	152.72	179.08	178.84	145.85	142.24	141.68	128.68	83.83	110.77	*	*
HEMBA1001804	23.86	24.89	37.02	23.72	21.32	19.26	15.97	14.33	15.78	*	-
HEMBA1001808	43.55	20.96	68.71	87.7	43.83	57.55	24.58	18.07	23.43		
HEMBA1001815	6.42	4.63	6.88	6.61	4.72	4.09	3.11	2.02	10.18		
HEMBA1001820	30.69	48.68	53.49	25.04	17.81	22.87	7.35	8.47	10.42	*	**
HEMBA1001821	8.82	12.28	20.89	18.14	12.47	19.32	10.67	8.58	14.11		
HEMBA1001835	3.29	3.32	8.82	2.2	2.68	3.13	2.05	0.94	1.44		

7 / 157

【0831】

【表360】

(1228)

特開2002-191363

2453

2454

HENBA1001639	3.01	3.66	7.63	2.63	2.16	1.96	2	2.43	2.09				
HENBA1001640	29.8	42.06	50.77	23.27	24.57	21.38	34.37	24.57	21.57	*	-		
HENBA1001647	5.45	3.94	16.28	4.05	8.97	2.61	5.21	2.15	4.19				
HENBA1001651	40.26	49.2	70.61	64.68	55.88	51.2	30.59	17.33	15.12	*	-		
HENBA1001655	10.17	9.88	20.1	6.79	14.34	13.87	15	5.01	12.4				
HENBA1001658	5.48	21.34	12.68	3.21	27.79	21.94	17.32	4.91	4.48				
HENBA1001661	1.92	2.27	3.68	1.72	2.22	1.93	2.04	0.71	3.48				
HENBA1001665	5.27	5.54	9.55	3.17	2.96	2.48	2.94	1.87	2.02	*	*	-	
HENBA1001670	14.59	18.93	22.5	10.01	11.18	7.78	13.23	9.01	14.55	*	*	-	
HENBA1001672	5.67	6.19	14.07	4.38	5.16	2.99	5.11	2.78	4.53				
HENBA1001673	67.5	83.13	103.8	79.62	78.6	82.68	79.54	46.75	60.49				
HENBA1001675	11.65	14.39	26.02	11.92	12.54	11.44	8.96	6.24	12.74				
HENBA1001678	5.94	6.55	16.33	9.03	9.2	9.7	5.97	3.49	7.17				
HENBA1001678	408.84	480.84	545.99	212.4	345.11	245.37	243.82	358.42	324.52	*	*	-	
HENBA1001680	17.81	21.9	34.13	4.77	16.15	9.65	5.24	7.39	4.29	*	*	-	
HENBA1001681	4.07	7.09	7.53	4.19	6.03	3.95	5.56	3.95	9.27				
HENBA1001684	9.88	10.78	16.76	5.52	7.17	9.85	8.63	7.5	7.69				
HENBA1001695	6.3	10.78	12.26	4.69	5.88	5.01	6.15	3.07	8.59				
HENBA1001702	4.21	4.98	12.28	2.87	3.73	2.48	2.67	0.6	1.75				
HENBA1001709	2.14	4.28	13.24	2.65	2.4	2.49	3	0.8	2.49				
HENBA1001711	4.48	2.9	7.77	4.65	5.15	6.37	2.22	1.03	3.22				
HENBA1001712	17.3	21.73	24.38	4.68	4.43	7.93	6.13	3.94	7.49	**	**	-	
HENBA1001714	5.44	5.06	12.28	3.87	6.98	8.98	4.35	1.75	2.27				
HENBA1001717	12.02	13.24	20.14	10.04	12.58	8.85	8.97	5.33	8.75				
HENBA1001718	6.01	7.09	16.45	6.91	7.4	8.4	6.45	5.23	4.35				
HENBA1001723	4.38	5.84	13.91	3.89	3.74	3.51	3.82	2.35	1.25				
HENBA1001731	2.72	5.88	12.37	2.09	2.33	1.98	2.59	1.24	1.73				
HENBA1001734	3.48	4.88	15.18	4.34	3.76	3.13	3.75	1.03	2.79				
HENBA1001735	7.38	10.88	24.38	12.76	10.32	13.2	4.98	2.9	5.27				
HENBA1001741	15.06	22.98	34.81	18.57	17.99	31.53	18.54	14.22	21.03				
HENBA1001744	3.63	4.77	7.03	5.75	5.45	4.88	3.08	0.38	1.67				
HENBA1001745	6.61	5.99	11.87	4.17	5.23	3.45	2.59	2.9	1.77	*	-		
HENBA1001748	10	14.3	20.05	11.55	8.04	11.63	7.31	4.66	10.54				
HENBA1001761	4.09	4.47	16.61	2.95	5.47	4.97	4.55	2.51	3.77				
HENBA1001762	12.27	16.22	31.71	11.79	12.77	9.07	10.3	5.04	7.64				
HENBA1001781	15.21	11.25	22.91	6.13	9.66	7.74	12.65	6.92	10.68				
HENBA1001784	7.14	8.4	14.02	11.89	8.88	32.45	4.84	4.18	5.94				
HENBA1001791	5.49	5.04	11.37	7.91	5.03	6.97	4.96	3.16	6.54				
HENBA1001794	20.29	19.03	24.02	20.07	12.65	17.58	18.71	4.14	12.87				
HENBA1001800	3.21	4.6	6.87	7.34	5.09	5.03	5.14	1.64	4.39				
HENBA1001803	13.17	17	28.4	25.23	15.99	17.41	14	9.49	14.18				
HENBA1001804	10.28	12.82	18.21	18.52	13.37	6.78	5.08	3.75	9.33				
HENBA1001808	1.39	1.46	0.67	2.77	3.18	1.88	0.98	1.2	0.66	*	*	-	
HENBA1001809	10.03	14.07	29.16	27.08	14.98	19.9	10.22	6.78	12.88				
HENBA1001811	4.85	4.27	3.18	10.29	3.63	4.31	3.6	2.84	5.25				
HENBA1001815	12.35	11.27	24.93	14.76	10.57	16.04	11.75	6.51	11.06				
HENBA1001816	15.52	16.02	31.22	15.84	13.09	17.05	15.48	10.68	15.2				
HENBA1001819	6.04	14.01	18.43	8.04	9.05	17.69	7.85	3.75	9.93				
HENBA1001820	1.8	2.22	6.75	3.5	2.81	3.45	1.98	0.63	2.02				
HENBA1001822	10	12.84	23.21	14.57	16.12	12.64	12.36	8.85	15.78				
HENBA1001824	5.2	8.63	7.51	6.46	8.51	5.78	4.87	0.96	4.7				
HENBA1001835	9.56	9.62	14.11	4.64	5.39	6.11	3.44	4.66	3.04	*	**	-	
HENBA1001844	8.02	10.52	18.05	12.42	10.73	9.57	5.22	6.54	4.99				
HENBA1001847	5.48	6.03	7.01	5.38	4.93	4.88	3.96	3.19	3.69	**	-		
HENBA1001849	6.93	9.14	14.78	5.03	7.88	7.21	7.8	8.87	10.07				
HENBA1001850	16.31	17.09	25.61	16.6	20.16	20.58	11.51	7.47	10.97	*	-		
HENBA1001861	2.35	3.42	4.47	4.07	3.37	3.89	2	1.11	3.84				
HENBA1001862	13.65	31.74	17.1	16.22	16.82	16.11	10.11	2.6	10.67				
HENBA1001864	2.79	3.87	7.65	5.32	5.64	4.4	3.29	2.25	6.43				
HENBA1001866	5.67	7.28	17.71	11.53	8.34	6.82	3.42	4.81	5.52				
HENBA1001869	12.86	19.53	27.68	23.68	16.72	23.14	5.91	10.06	12.62				
HENBA1001871	3.55	6.35	6.61	4.54	2.2	3.49	3.21	1.17	2.75				
HENBA1001876	4.6	3.93	8.22	5.18	4.46	4.39	3.78	4.15	6.11				
HENBA1001878	8.8	14.72	27.92	7.18	13.52	10.62	6.43	4.37	4.82				

8/157

【0832】

【表361】

2455

2456

MEMBA1001879	23.8	24.57	36.17	27.25	30.71	21.12	18.92	12.59	21.51										
MEMBA1001884	22.79	25.41	39.57	18.38	21.7	20.15	24.74	18.18	27.02										
MEMBA1001886	2.49	3.21	10.85	4.21	4.2	3.88	3.99	2.94	2.81										
MEMBA1001888	5.93	5.79	13.13	8.79	5.21	8.71	5.32	6.52	7.0										
MEMBA1001890	12.55	17.2	16.26	7.93	7.84	9.81	5.34	7.04	9.79	*	*								
MEMBA1001898	8.62	13.4	18.92	8.94	9.97	13.19	7.37	7.12	11.57										
MEMBA1001899	9.63	9.32	14.85	9.07	9.12	9.82	4.74	3.78	7.78										
MEMBA1001904	17.69	9.43	25.81	8.49	15.82	16.23	7.42	3.21	4										
MEMBA1001910	12.84	17.03	18.88	10.76	16.22	11.5	15.72	11.44	11.3										
MEMBA1001911	84.18	139.96	104.54	60.48	92.99	38.82	38.97	44.93	16.51	*									
MEMBA1001912	8.8	18.15	15.1	11.38	13.1	11.76	6.02	5.45	5.48	*									
MEMBA1001913	63.6	87.85	178.77	91.87	76.48	71.89	38.1	40.9	78.58										
MEMBA1001915	9.95	12.39	24.42	17.53	14.5	13	8.17	4.57	10.88										
MEMBA1001918	5.57	6.48	12.5	8.11	6.81	8.09	4.3	2.18	7.19										
MEMBA1001921	8.68	9.91	12.62	8.57	8.25	7.86	5.28	3.66	4.78	**									
MEMBA1001931	9.24	10.47	21.38	7.16	7.68	8.88	8.41	8.87	7.96										
MEMBA1001939	3.59	7.32	15.17	1.26	16.55	6.08	4.84	5.03	14.82										
MEMBA1001940	11.2	9.93	21.45	9.87	12.2	10.55	7.02	5.09	7.03										
MEMBA1001942	22.25	38.8	27.34	8.22	35.32	31.27	17.28	9.8	12.44	*									
MEMBA1001944	14.23	17.39	30.07	21.25	14.66	13.47	4.56	8.32	14.97										
MEMBA1001945	8.53	9.04	18.07	9.58	8.68	6.81	6.7	4.7	10.21										
MEMBA1001950	35.24	35.06	25.81	22.39	20.56	21.78	31.55	22.8	14.77	*									
MEMBA1001951	39.37	38.17	52.87	44.58	33.39	34.05	41.89	29.81	45.03										
MEMBA1001958	3.68	3.88	9.18	1.87	2.36	1.68	2.38	1.43	3.59										
MEMBA1001960	11.92	13.79	24.84	7.91	13.38	11.06	7.97	3.58	9.09										
MEMBA1001962	10.62	11.31	28.75	9.3	13.78	9.37	10.95	7.24	13.69										
MEMBA1001964	11.99	14.1	28.18	9.47	13.95	12.22	12.21	4.77	10.5										
MEMBA1001967	33.18	31.11	63.39	22.3	13.19	20.77	8.29	10.21	24.27										
MEMBA1001979	6.77	6.74	6.61	3.38	4.61	6.79	4.27	2.37	4.75	*									
MEMBA1001987	6.84	6.64	7.7	5.03	3.79	3.32	4.73	5.25	2.63	*									
MEMBA1001991	4.14	2.8	6.6	1.78	2.7	1.18	2.19	1.73	1.57										
MEMBA1002003	3.19	3.64	8.31	2.58	2.51	1.13	4.14	2.93	2.53										
MEMBA1002005	21.43	25.67	43.84	22.88	20.88	16.44	24.78	11.27	20.55										
MEMBA1002008	15.2	19.08	39.8	14.05	18.61	13.82	20.77	11.98	21.03										
MEMBA1002018	4.58	11.59	12.53	2.51	5.88	20.49	4.43	1.81	2.74										
MEMBA1002022	28.07	36.45	75.11	52.83	42.93	48.33	30.25	25.12	55.78										
MEMBA1002029	19.99	24.25	30.34	10.7	17.32	16.55	10.55	13.15	11.58	*									
MEMBA1002030	11.08	12.89	19.22	8.24	7.3	5.76	8	6.82	6.24	*	*								
MEMBA1002035	3.07	17.04	7.87	2.25	5.11	2.18	4.84	3.89	6.08										
MEMBA1002037	95.17	141.19	140.82	59.83	84.82	73.74	61.45	42.2	28.14	*	*								
MEMBA1002038	11.98	19.11	18.49	5.44	8.02	6.74	4.83	2.27	2.59	*	**								
MEMBA1002039	8.42	7.41	13.49	2.37	5.91	2.48	3.27	1.37	3.18	*									
MEMBA1002042	3.25	8.61	11.52	6.6	8.85	6.58	7.97	0.75	4.67										
MEMBA1002043	13.24	16.58	22.47	5.13	8.87	8.08	5.01	5.72	8.43	*	*								
MEMBA1002048	25.65	31.58	36.48	5.14	12.63	17.17	11.27	12.56	24.5	*	*								
MEMBA1002049	431.22	564.03	858.02	234.02	525.06	482.28	304.41	585.28	233.29										
MEMBA1002053	68.35	81.58	114.43	104.42	98.85	104.79	58.5	40.85	89.44										
MEMBA1002055	18.16	20.77	33.11	22.38	16.8	17.16	15.07	11.5	15.77										
MEMBA1002058	5.28	6.55	10.95	2.52	2.93	4.72	5.3	2.16	5.03										
MEMBA1002061	3.02	4.19	8.85	2.45	3.45	0.92	2.37	0	1.82										
MEMBA1002080	2.59	8.98	20.11	3.27	8.34	5.7	8.09	2.75	4.82										
MEMBA1002084	7.01	3.85	10.52	5.98	3.47	4.97	2.89	1.31	2.02										
MEMBA1002085	3.68	5.88	10.92	2.9	5.24	7.31	3.65	1.95	5										
MEMBA1002092	18.94	17.78	38.17	14.49	14.47	17.01	14.19	7.25	10.23										
MEMBA1002098	8.74	16.29	15.2	3.59	8.82	8.02	9.31	14.32	5.83										
MEMBA1002100	13.69	10.18	19.02	11.56	6.49	4.52	8.34	7.69	7.11										
MEMBA1002101	12.82	14.09	31.83	14.79	12.78	11.05	18.17	8.88	13.98										
MEMBA1002102	4.18	7.54	13.28	4.1	4.71	4.58	3.77	3.14	3.09										
MEMBA1002105	5.07	7.15	15.22	5.77	3.71	9.62	4.21	1.98	2.79										
MEMBA1002107	7.22	7.4	21.27	8.53	10.18	12.18	2.67	2.93	5.58										
MEMBA1002113	5.14	7.17	14.68	10.73	7.63	8.58	8.5	2.35	7.99										
MEMBA1002119	24.1	28.98	60	38.03	31.14	28.41	31.91	16.47	21.72										
MEMBA1002125	4.31	5.62	8.68	11.65	3.83	6.15	2.02	2.4	2.97										
MEMBA1002131	2812	1876.5	3969.7	1308.5	2879.6	1519.6	1255.8	1938.4	1851.4										

9 / 157

【0833】

【表362】

2457

2458

HEMBA1002133	28.03	35.39	76.13	107.6	70.61	84.67	28.03	14.5	31.23
HEMBA1002139	11.59	23.7	52.64	28.04	21.84	20.31	32.1	12.89	25.76
HEMBA1002141	3.72	5.77	10.55	5.21	5.75	3.18	3.83	1.34	5.31
HEMBA1002144	2.85	4.6	6.42	4.94	2.2	5.21	1.49	2.58	2.66
HEMBA1002147	10.27	15.44	23.81	11.25	8.38	13.35	8.76	6.54	9.55
HEMBA1002150	1.33	1.23	3.53	2.48	2.28	2.8	1.93	0.73	2.03
HEMBA1002151	6.03	4.63	11.41	5.99	5.58	6.88	5.34	2.78	4.91
HEMBA1002153	0.77	2.83	4.03	3.48	4.19	3.78	2.71	1.39	3.61
HEMBA1002156	2.06	3.91	4.83	5.6	4.46	2.69	2.87	0.76	2.49
HEMBA1002160	58.44	32.19	84.15	108.2	70.78	78.85	43.2	16.4	44.98
HEMBA1002161	44.7	21.27	41.95	81.85	58.89	51.89	26.74	12.83	24.21
HEMBA1002162	89.86	103.48	226.88	313.25	143.99	221.78	91.22	73.86	98.69
HEMBA1002183	40.99	33.35	79.59	79.2	53.49	69.59	30.85	22.17	36.21
HEMBA1002184	5.75	8.38	13.54	8.04	3.83	5.52	3.45	2.03	3.62
HEMBA1002186	15.02	15.38	25.99	20.34	17.99	15.75	15	11.24	18.28
HEMBA1002187	11.24	16.13	23.3	10.71	13.78	13.32	12.46	7.17	15.43
HEMBA1002173	2.74	4.74	7.08	7.8	4.98	8.28	4.32	0.11	6.88
HEMBA1002177	3.19	7.92	6.73	9.48	13.68	11.32	3.6	0.89	5.89
HEMBA1002178	3.07	3.14	5.24	4.35	5.65	4.54	2.7	1.84	3.18
HEMBA1002179	4.9	4.72	8.91	8.85	4.93	4.63	2.62	2.88	3.55
HEMBA1002185	24.67	32.27	51.89	14.08	21.43	18.77	8.38	11.95	12.92
HEMBA1002188	5.88	6.14	7.11	2.87	4.39	3.33	2.27	4.54	2.88
HEMBA1002189	69.59	81.69	157.25	233.85	141.87	162.77	75.89	48.73	66.24
HEMBA1002191	4.08	3	7.85	6.05	7	4.24	3.01	1.76	4.2
HEMBA1002192	5.95	7	11.91	7.01	8.79	8.52	4.84	2.39	5.33
HEMBA1002195	4.48	6.6	10.03	6.4	8.23	6.42	6.4	2.77	4.74
HEMBA1002196	11.48	14.42	16.69	14.62	12.58	12.05	7.85	1.78	5.08
HEMBA1002199	17.19	29.45	44.34	8.12	15.7	10.5	4.22	6.66	8.91
HEMBA1002204	12.38	17.29	25.35	18.54	15.85	13.06	6.95	7.55	12.67
HEMBA1002208	4.42	3.1	8.67	2.78	2.3	3.81	1.87	2.17	2.61
HEMBA1002212	8.54	9.41	18.06	6.8	5.49	9.18	8.07	6.31	7.72
HEMBA1002215	3.77	4.75	11.1	3.82	3.84	4.93	3.19	2.31	3.82
HEMBA1002217	3.21	2.97	8.05	1.88	2.5	3.28	2.77	1.7	3.42
HEMBA1002220	40.07	58.29	38.41	33.44	37.1	28.5	27.12	21.87	28.71
HEMBA1002226	3.85	8.37	9.01	6.94	8.12	6.13	6.11	1.84	2.28
HEMBA1002227	29.2	48.92	80.06	21.28	24.93	28.49	9.08	21.04	35.76
HEMBA1002229	32.53	65.16	51.1	12.99	38.46	38.19	10.92	21.02	13.32
HEMBA1002237	6.24	7.25	13.43	6.25	5.49	6.14	5.27	3.9	7.13
HEMBA1002239	4.63	2.53	7.81	2.56	2.45	2.52	2.39	1.83	5.54
HEMBA1002241	108.58	86.21	101.41	181.82	99.49	118.15	85.67	62.74	65.24
HEMBA1002253	12.22	17.16	20.51	2.71	7.33	6.42	3.87	2.39	2.33
HEMBA1002257	7.54	10.24	15.29	4.58	8.87	7.35	4.9	4.18	3.85
HEMBA1002259	13.5	22.26	29.8	14.68	19.62	15.23	17.99	11.27	18.91
HEMBA1002262	15.09	23.24	29.3	18.72	15.42	11.55	9.03	9.95	14.69
HEMBA1002265	14.18	19.86	21.44	12.15	13.48	12.48	8.23	10.18	13.41
HEMBA1002267	27.68	15.47	60.07	10.16	19.76	26.78	8.61	9.81	8.8
HEMBA1002270	6.74	5	13	3.57	4.78	3.79	3.08	1.88	3.1
HEMBA1002286	3.25	3.45	8.41	1.82	1.88	2.17	2.49	0.44	3.13
HEMBA1002290	13.68	14.35	29.79	17.54	14.92	16.25	8.88	7.54	11.64
HEMBA1002302	2.59	3.21	9.4	1.67	2.25	2.2	2.59	2.03	2.91
HEMBA1002304	28.75	42.22	44.71	47.37	56.93	39.52	32.89	19.02	22.8
HEMBA1002307	11.61	12.1	29.02	16.8	17.62	14.8	7.63	7.75	11.14
HEMBA1002316	59.84	53.59	78.81	100.13	70.41	88.41	40.44	34.11	39.43
HEMBA1002319	5.85	5.03	12.11	8.62	5.15	5.86	4.45	4.18	4.84
HEMBA1002320	5.15	6.41	11.19	4.43	3.25	5.88	2.72	1.07	1.82
HEMBA1002321	28.14	32.72	45.81	15.38	15.91	12.15	25.74	14.29	26.13
HEMBA1002328	3.49	3.33	9.66	1.53	2.84	1.89	1.87	1.15	3.61
HEMBA1002333	14.03	17.53	33.87	10.89	17.53	10.54	14.85	12.2	18.17
HEMBA1002337	74.1	90.11	104.84	39.01	88.82	67.57	65.51	46.31	39.48
HEMBA1002339	7.89	5.95	12.61	11.75	8.3	7.98	3.81	2.17	4.39
HEMBA1002341	10.07	14.65	15.74	8.32	6.89	8.79	4.57	2.09	3.78
HEMBA1002348	59.9	42.49	67.67	22.81	39.28	47.94	42.75	47.55	14.2
HEMBA1002349	8.79	4.77	8.57	3.9	2.41	2.17	3.1	1.47	1.4
HEMBA1002353	16.84	19.17	38.78	16.44	19.81	18.38	31.34	14.04	19.17

10 / 157

【0834】

【表363】

2460

HEBIA1002356	38	42.3	70.07	20.6	28.38	26.6	38.92	22.92	28.48
HEBIA1002357	5.52	13.1	13.45	8.21	11.98	5.72	6.55	2.43	8.78
HEBIA1002360	7.18	14.99	31.64	17.74	12.78	10.39	10.08	6.73	9.23
HEBIA1002363	7.21	8.56	15.23	7.32	10.07	12.95	6.94	4.08	5.51
HEBIA1002365	6.13	6.37	6.27	3.08	3.21	2.92	4.54	1.5	1.04
HEBIA1002370	2.68	3.2	4.12	1.48	1.53	1.16	1.82	1.08	0.59
HEBIA1002374	7.68	8.73	12.15	4.74	6.3	4.18	4.71	3.04	1.54
HEBIA1002378	3.93	7.64	11.22	5.3	6.08	2.77	7.34	1	2.6
HEBIA1002377	0.99	9.89	16.86	5.63	7.09	6.16	6.12	1.21	4.87
HEBIA1002380	16.33	16.4	28.08	14.11	13.73	11.4	15.31	7.9	14.89
HEBIA1002381	4.06	9.29	16.58	7.37	8.4	6.25	6.66	1.71	0.19
HEBIA1002384	8.47	11.56	16.49	7.98	8.37	10.32	10.3	5.34	12.08
HEBIA1002388	11.42	20.29	29.05	6.75	14.68	8.45	8.49	15.83	6.08
HEBIA1002396	12.68	16.26	27.27	7.41	9.5	8.56	5.14	3.17	3.81
HEBIA1002402	12.79	14.83	21.89	12.15	10.97	7.11	8.38	8.96	10.94
HEBIA1002417	18.84	19.38	28.26	14.12	13.8	10.71	18.74	11.92	12.94
HEBIA1002419	5.78	8.38	11.39	2.24	4.5	1.91	3.8	1.74	2.93
HEBIA1002420	19.24	23.07	61.09	28.34	25.11	18.31	20.11	12.19	18.92
HEBIA1002421	18.39	32.78	48.68	28.38	38.24	25	28.06	7.94	14.32
HEBIA1002423	22.61	32.68	44.05	7.08	7.68	6.56	4.68	5.6	8.16
HEBIA1002424	18.87	10.88	26.72	15.1	10.87	9.51	12.1	6.14	11.48
HEBIA1002426	7.39	16.83	33.68	10.19	19.75	11.23	14.09	9.38	9.51
HEBIA1002430	3.4	2.89	7.25	1.05	6.01	3.49	1.89	3.82	1.55
HEBIA1002439	16.88	14.68	38.41	6.15	12.09	6.57	10.43	3.04	5.71
HEBIA1002441	6.85	11.02	20	8.08	6.54	5.94	8.38	4.52	8.82
HEBIA1002454	12.67	10.57	41.83	18.84	16.12	15.91	15.68	6.28	10.15
HEBIA1002458	16.76	29.12	42.05	23.99	29.68	19.78	21.68	9.8	15.93
HEBIA1002460	10.75	22.65	60.16	30.55	31.38	38.17	25.74	12.29	26.18
HEBIA1002462	4.28	4.96	6.87	4.04	2.62	5.54	6.09	4.68	1.34
HEBIA1002465	18.48	17.14	33.42	10.69	9.15	14.44	4.41	4.26	2.08
HEBIA1002469	5.68	10.77	19	7.83	8.55	8.09	9.48	5.81	6.88
HEBIA1002475	6.8	4.95	7.87	3.63	3.61	2.62	2.18	0.88	3.68
HEBIA1002477	14.5	19.14	47.19	10.8	18.29	17.91	11.51	8.53	9.54
HEBIA1002480	3.04	4.81	11.31	4.98	4.91	2.26	3.45	1.28	2.5
HEBIA1002481	3.73	5.8	10.38	6.88	9.38	9.35	6.55	0.82	5.3
HEBIA1002486	14.96	20.73	38.85	17.9	13.84	17.9	9.8	9.17	15.34
HEBIA1002490	32.89	43.68	75.38	48.87	29.68	38.22	19.18	16.64	20.65
HEBIA1002495	43.66	58.23	82.65	111.1	32.51	41.26	76.83	29.1	14.68
HEBIA1002498	3.78	6.46	9.54	6.71	6.27	12.7	10.64	3.53	6.2
HEBIA1002501	7.77	5.28	15.06	10.61	8.44	10.8	6.68	2.78	7.39
HEBIA1002503	1.3	3.78	4.26	12.14	4.16	3.06	2.94	0.74	2.48
HEBIA1002504	11.32	6.35	15.09	5.12	7.76	8.01	3.02	1.3	4
HEBIA1002508	0.86	9.32	4.01	3.71	7.02	6.17	4.27	2.16	3.39
HEBIA1002513	9.71	15.43	17.71	11.5	8.94	8.42	3.65	5.7	5.29
HEBIA1002515	22.68	29.58	33.68	19.66	17.58	7.29	3.31	4.58	5.25
HEBIA1002524	2.62	5.5	6.81	5.24	5.07	5.47	3.51	2.04	6.31
HEBIA1002538	4.68	8.27	12.8	5.54	8.68	8.78	7.78	3.44	3.38
HEBIA1002542	14.28	22.48	34.17	28.25	22.99	20.23	26.21	13.64	24.14
HEBIA1002544	80.29	71.81	142.33	181.63	103.72	120.69	75.35	42.62	73.65
HEBIA1002546	16.48	22.78	27.12	14.78	16.43	17.06	9.62	3.52	9.4
HEBIA1002547	3.95	4.98	8.23	6.73	9.06	8.15	9.5	2.38	7.03
HEBIA1002550	49.15	78.71	80.52	40.75	28.92	26.29	11.49	29.11	24.84
HEBIA1002551	21.38	26.41	43.07	27.63	15.88	20.68	15.81	13.41	19.29
HEBIA1002552	2.51	5.57	9.92	2.51	4.28	4.71	1.72	3.19	4.1
HEBIA1002555	22.2	20.07	42.05	23.18	22.23	22.78	15.94	12.91	13.98
HEBIA1002568	12.08	15.97	19.47	12.44	11.51	11.85	16.99	12.78	19.34
HEBIA1002581	1.12	5.08	10.95	9.63	4.84	7.75	3.52	2.9	5.93
HEBIA1002582	5.102	16.11	24.74	19.28	28.78	27.58	31.51	7.07	19.83
HEBIA1002586	196.94	317.85	248.69	300.72	297.51	250.18	199.37	80.64	130.04
HEBIA1002589	28.5	41.31	68.88	12.64	22.81	12.48	7.29	12.02	18.03
HEBIA1002570	30.8	40.14	52.73	26.65	26.92	32.98	18.03	21.81	22.78
HEBIA1002574	18.81	26.28	28.64	14.11	18.09	18.3	12.08	11.37	15.9
HEBIA1002583	7.94	6.71	16.38	3.31	5.66	5.07	2.73	4.44	3.12
HEBIA1002587	2.96	4.41	12.22	7.72	5.53	9.69	6.2	2.49	5.6

11 / 157

2461

HENBA10025901	5.95	11.3	17.91	3.88	9.68	7.63	4.92	3.2	3.29
HENBA1002592	24.58	35.63	62.38	31.85	31.39	28.3	31.44	17.58	25.5
HENBA1002595	18.13	24.94	28.1	20.87	22.49	18.85	23.31	9.38	10.88
HENBA1002608	10.82	12.83	20.23	14.42	14.83	14.25	5.32	5.93	14.35
HENBA1002617	4.27	5.73	7.4	1.57	2.4	3.96	1.42	3.15	4.02
HENBA1002619	2.86	2.62	7.5	3.1	2.95	3.67	1.84	0.73	2.61
HENBA1002621	8.59	8.17	23.61	10.11	9.98	16.3	7.86	5.98	9.09
HENBA1002624	27.35	35.73	38.2	18.75	33.78	30.42	15.63	13.42	13.94
HENBA1002628	9.91	18.29	22.67	1.48	9.73	6.58	2.84	2.88	1.66
HENBA1002629	17.5	28.89	38.07	15.14	20.64	15.47	19.33	12.69	24.63
HENBA1002632	12.9	25.94	21.19	10.8	16.54	18.64	7.96	5.15	6.22
HENBA1002645	18.16	17.89	28.67	12.29	14.08	14.33	9.35	8.1	11.98
HENBA1002651	13.28	18.44	26.72	15.43	10.74	15.28	12.68	8.84	17.98
HENBA1002652	39.73	48.42	65.63	48.16	37.25	33.78	28.28	20.1	35.38
HENBA1002659	9.03	11.88	19.15	8.75	10.05	12.88	11.02	6.8	9.75
HENBA1002661	5.06	3.99	9.25	2.67	2.42	2.71	2.5	0.54	3.81
HENBA1002666	4.98	2.76	10.31	1.02	2.15	2.47	1.85	1.39	2.97
HENBA1002687	88.28	92.17	95.25	40.2	48.75	38.84	17.08	15.46	18.87
HENBA1002673	18.24	11.61	24.1	10.91	20.39	11.78	14.29	5.19	10.4
HENBA1002678	13.62	18.9	33.62	20.45	17.4	13.57	11.48	10.78	12.98
HENBA1002679	13.69	14.71	16.04	4.7	9.66	9.85	3.39	2.88	4.57
HENBA1002688	5.8	5.12	9.48	4.59	5.44	5.04	4.89	4.1	8.84
HENBA1002696	59.06	73.44	43.41	32.21	38.98	39.85	22.83	22.78	43.71
HENBA1002703	7.17	12.58	16.11	7.07	6.09	6.51	6.44	2.28	5.09
HENBA1002706	6.13	6.22	16.32	2.59	4.89	3.24	4.44	2.11	3.85
HENBA1002712	11.08	16.22	21.38	7.53	8.92	6.72	6.23	2.15	5.09
HENBA1002715	7.25	9.24	15.83	3.34	13.16	4.85	8.59	3.19	5.48
HENBA1002716	10.24	5.79	8.57	7.72	6.63	11.75	6.68	8.82	8.25
HENBA1002718	5.54	9.62	8.18	5.1	4.05	3.87	4.38	2.74	1.77
HENBA1002728	12.44	14.25	13.51	6.78	9.73	5.64	5.16	3.78	4.32
HENBA1002730	18.89	21.85	36.97	19.48	14.69	13.58	21.24	13.64	21.33
HENBA1002734	18.2	30.15	26.54	25.61	18.78	20.98	17.72	11.44	17.14
HENBA1002742	3.87	2.34	11.09	2.03	2.88	2.81	1.76	0.66	1.41
HENBA1002746	22.55	32.65	52.23	26.89	25.86	22.2	22.76	13.04	18.94
HENBA1002749	9.65	15.62	22.08	8.63	12.94	6.47	8.84	2.48	4.44
HENBA1002750	4.68	4.85	9.25	7.27	4.67	4.79	3.27	2.62	3.14
HENBA1002755	6.97	7.74	9.25	6.37	7.77	4.18	3.95	4.22	4.38
HENBA1002759	16.7	17.27	27.62	16.79	11.68	13.5	9.91	9.05	10.68
HENBA1002763	47.58	57.94	58.78	64.23	38.74	38.37	38.64	19.12	21.85
HENBA1002767	8.72	14.41	17.03	5.04	11.17	4.98	8.31	3.05	8.44
HENBA1002768	5.62	5.94	12.79	2.88	4.3	2.49	2.51	1.17	1.65
HENBA1002769	17.19	27.21	15.65	8.55	10.93	3.03	6.22	1.99	5.25
HENBA1002770	2.13	4.12	10.85	3.83	4.73	3.88	2.44	1.5	3.81
HENBA1002777	6.28	5.97	10	5.84	6.03	4.58	3.3	3.36	7.09
HENBA1002779	4.32	3.05	9.05	3.43	3.09	3.54	2.97	2.09	4.26
HENBA1002780	18.15	7.73	29.43	3.93	22.04	8.61	11.74	14.31	4.26
HENBA1002790	44.22	50.48	73.09	31.77	35.83	33.38	43.74	34.78	30.04
HENBA1002794	7.76	9.01	14.22	4.88	8.33	4.1	3.98	2.38	3.09
HENBA1002798	58.18	53	86.62	94.91	76.85	71.63	49.35	25.85	43.83
HENBA1002801	3.51	8.44	18.4	3.59	4.68	6.72	3.9	1.17	4.1
HENBA1002810	3.05	9.67	16.69	8.92	5.81	6.43	5.04	3.55	5.63
HENBA1002816	33.64	55.39	45.67	6.42	16.08	37.06	10.38	10.74	17.79
HENBA1002818	13.01	14.09	20.64	7.78	6.78	5.49	5.04	2.98	3.82
HENBA1002820	27.01	29.21	56.68	35.9	30.38	30.6	30.02	21.79	27.81
HENBA1002826	19.2	24.63	59.75	28.13	24.2	21.9	24.8	13.89	24.13
HENBA1002833	63.26	64.77	137.72	30.62	26.08	40.67	37.5	33.73	24.96
HENBA1002850	32.24	50.44	106.87	13.72	14.84	20.17	21.65	20.43	7.04
HENBA1002852	2.91	4.02	13.2	2.77	4.61	2.19	2.67	1.11	1.62
HENBA1002853	5.54	4.99	10.17	2.83	5.88	5.71	5.41	0.6	2.81
HENBA1002867	12.64	16.67	40.23	22.88	17.48	25	12.79	7.67	12.96
HENBA1002876	33.33	40.68	58.24	16.95	25.21	41.71	19.38	18.11	7.35
HENBA1002886	38.07	46.59	100.23	134.31	130.1	87.32	31.47	23.92	40.5
HENBA1002896	379.7	87.77	442.56	153.27	289.24	233.48	153.73	231.71	129.4
HENBA1002913	5.91	13.05	17.5	7.91	16.67	18.64	16.24	3.53	12.2

12 / 157

2462

2462

2464

HEBHA1002921	10.68	17.69	28.78	4.57	11.62	17.13	10.57	6.89	8.38
HEBHA1002924	16.62	23.08	41.44	17.55	21.51	18.61	18.89	6.29	12.01
HEBHA1002934	7.43	7.93	16.24	8.33	10.71	10.41	9.2	4.4	7.82
HEBHA1002935	44.38	58.85	52.49	18.89	32.69	26.1	15.32	13.88	12.84
HEBHA1002937	2.13	5.1	6.49	6.16	3.25	2.49	3.22	1.79	2.89
HEBHA1002939	6.4	9.23	9.0	6.39	7.51	3.18	2.69	1.8	1.63
HEBHA1002944	8.74	7.31	12.93	7.18	6.53	7.34	7.98	2.7	5.69
HEBHA1002951	2.28	3.34	4.29	3.26	3.2	3.99	3	1.79	2.41
HEBHA1002954	17.42	19.44	25.7	12.87	16.68	13.48	13.78	7.82	14.9
HEBHA1002982	8.78	11.09	12.46	11.77	11.02	6.43	11.04	6.23	10.79
HEBHA1002983	6.03	12.43	15.52	4.99	4.92	4.17	4.83	4.54	3.33
HEBHA1002970	3.97	3.94	8.85	9.45	5.39	4.2	3.04	1.94	3.36
HEBHA1002971	45.35	66.82	65.65	55.83	32.32	29.43	18.21	22.18	18.43
HEBHA1002973	9.1	22.68	32.02	18.54	14.79	14.79	14.19	10.32	16.02
HEBHA1002978	16.61	14.49	12.58	8.49	10.35	10.49	8.97	8.63	9.32
HEBHA1002981	7.62	7.65	10.48	7.25	7.54	7.91	5.93	4.75	6.33
HEBHA1002985	117.94	141.28	115.79	100.57	91.73	67.47	38.07	32.99	67.85
HEBHA1002986	1.09	4.19	2.41	1.9	4.92	2.49	2.41	1.95	2.68
HEBHA1002988	6.78	12.45	10.78	6.53	13.58	5.88	8.35	4.68	5.68
HEBHA1002992	5.15	10.06	11.86	10.23	8.88	4.79	4.31	25.69	8.31
HEBHA1002995	4.3	8.89	7.73	5.51	4.65	4.85	3.59	3.01	4.66
HEBHA1002997	107.63	114.87	183.54	177.35	158.59	181.54	82.75	91.38	112.69
HEBHA1002999	53.34	68.01	72.24	37.38	47.87	41.25	62.98	43.59	63.81
HEBHA1003004	6.44	4.33	3.72	2.58	3.87	3.85	3	1.67	3.06
HEBHA1003006	4.28	6.06	7.7	1.79	6.77	4.05	4.37	2.61	3.52
HEBHA1003008	27.15	36.55	38.8	28.92	32.63	23.56	33.58	25.15	30.76
HEBHA1003021	5.81	12.5	13.45	11.13	6.12	5.84	7.32	6.94	6.02
HEBHA1003027	16.79	28.5	31.88	21.26	18.71	19.18	18.11	14.67	19.02
HEBHA1003029	19.01	29.05	31.25	23.38	17.83	18.43	15.34	15.3	18.09
HEBHA1003031	23.27	32.76	35.62	18.12	20.06	19.05	22.67	22.89	27.58
HEBHA1003032	10.61	18.48	23.67	10.46	10.87	8.83	13.71	10.94	13.4
HEBHA1003033	3.28	3.5	9.21	2.58	2.89	3.45	2.32	2.32	3.69
HEBHA1003034	32.22	43.17	44.48	25.3	23.69	20.88	30.34	23.98	27.83
HEBHA1003035	6.89	6.34	8.49	7.01	5.89	4.57	6.73	1.95	2.1
HEBHA1003037	3.15	4.70	8.43	8.95	8.5	6.33	1.17	2.19	4.48
HEBHA1003041	6.81	4.77	6.88	6.48	7.11	3.98	3.65	3.49	6.68
HEBHA1003046	5.59	5.67	7.87	2.55	3.97	2.74	1.89	1	3.82
HEBHA1003047	2.76	3.8	7.28	2.24	3.39	2.22	1.7	2.72	3.44
HEBHA1003048	7.12	9.91	17.38	6.18	6.88	5.24	7.18	5.24	6.81
HEBHA1003084	7.42	7.15	11.39	8.45	5.88	2.85	4.01	3.53	6.28
HEBHA1003087	3.5	4.14	10.69	1.88	3.75	2.7	1.84	2.09	2.56
HEBHA1003071	4.3	6.5	8.89	2.07	3.4	1.57	3.49	0.09	2.65
HEBHA1003072	22.65	30.56	72.87	40.85	32.84	38.51	17.84	22.4	33.19
HEBHA1003076	10.41	7.13	15.24	12.87	5.59	14.57	9.56	5.38	7.99
HEBHA1003077	67.77	42.58	38.04	15.22	22.48	42.9	9.06	21.41	25.7
HEBHA1003078	5.9	3.99	8.44	6.05	5.47	3.08	3.19	1.13	4
HEBHA1003079	2.67	1.28	11.95	1.19	1.44	0	0.126	1.15	2.6
HEBHA1003083	17.29	19.57	26.48	12.28	18.41	11.4	17.95	11.77	14.23
HEBHA1003086	68.37	81.42	76.05	81.77	61.08	81.6	45.88	46.71	41.69
HEBHA1003090	27.68	31.13	33.36	17.19	38.88	20.27	13.44	24.06	7.13
HEBHA1003094	7.82	15.14	35.58	19.45	14.45	10.58	8.94	6.32	19.5
HEBHA1003096	5.15	5.97	7.89	4.67	5.25	3.62	2.44	1.23	5.08
HEBHA1003098	61.24	66.94	73.83	44.19	37.83	35.41	49.21	42.98	49.32
HEBHA1003101	55.72	62.33	39.6	28.21	27.13	38.6	19.35	33.07	15.18
HEBHA1003108	21.19	29.16	38.82	18.38	20.48	14.85	17.89	17.38	21.07
HEBHA1003114	18.2	22.18	27.85	11.32	13.05	11.32	15.03	12.24	12.98
HEBHA1003117	4.65	5.29	16.23	3.16	8.98	2.08	3.65	1.47	2.05
HEBHA1003120	28.44	23.85	62.74	11.15	27.45	17.66	13.62	6.31	12.11
HEBHA1003129	13.18	14.01	38.23	23.02	18.83	17.41	9	7.3	18.68
HEBHA1003133	18.25	22.53	34.42	24.65	19.61	15.34	21.84	16.17	21.18
HEBHA1003138	16.39	19.51	25.59	18.11	18.18	15.75	14.64	13.22	14.96
HEBHA1003142	14.58	16.27	15.48	9.65	8.16	8.78	8.14	5.28	9.82
HEBHA1003148	18.81	26.84	28.42	18.45	15.92	11.14	21.43	11.51	11.69
HEBHA1003151	8.82	10.43	18.65	5.42	8.75	3.58	10.39	7.34	8.14

13 / 157

2465

HERBA1003152	4.26	6.9	16.64	3.43	9.18	3.49	6.82	2.76	2.37
HERBA1003157	16.47	20.65	36.14	14.57	15.21	12.8	18.89	12.37	10.25
HERBA1003166	6.07	8.14	15.45	9.26	7.78	7.15	6.88	3	7.14
HERBA1003171	4.76	6.01	1.83	0.22	9.4	0.95	4.99	3.3	4.74
HERBA1003175	9.08	9.38	15.39	5.94	5.41	4.09	8.31	5.64	6.78
HERBA1003179	11.74	19.14	23.08	11.43	19.69	18.88	15.24	11.13	8.75
HERBA1003186	32.34	34.21	53.13	30.02	30.6	23.62	32.72	27.67	23.79
HERBA1003196	4.93	6.75	11.29	1.52	4.36	1.91	2.86	1.86	1.27
HERBA1003197	11.79	18.34	28.33	10.88	17.37	11.43	13.65	9.71	13.58
HERBA1003199	23.1	49.93	40.51	25.63	28.94	20.25	30.98	8.01	23.38
HERBA1003202	23.9	28.7	33.49	4.92	13.85	18.67	6.88	7.48	15.28
HERBA1003204	4.47	10.66	14.07	5.38	6.97	3.56	4.1	2.93	6.45
HERBA1003210	26.93	30.85	39.03	16.12	18.87	10.78	20.75	18.2	19.07
HERBA1003212	25.47	38.89	40.06	19.34	20.83	15.78	34.72	24.84	21.07
HERBA1003218	4.32	4.85	11.19	2.63	4.68	1.07	3.74	6.58	2.3
HERBA1003220	35.02	48.9	66.81	20.36	27.56	20.54	37.25	33.69	30.02
HERBA1003222	6.38	11.47	27.67	6.14	9.88	5.65	9.63	5.71	8.41
HERBA1003225	1.89	5.89	11.4	3.87	2.88	2	2.78	0.38	1.82
HERBA1003229	4.11	5.35	10.63	5.08	8.91	5.05	3.62	3.37	4.28
HERBA1003230	19	22.89	41.86	21.05	18.18	18.78	15.9	8.88	15.64
HERBA1003235	8.36	8.82	18.89	3.6	5.78	3.31	5.14	2.94	2.76
HERBA1003236	28.57	38.02	51.11	22.25	24.92	17.63	25.41	23	19.94
HERBA1003250	3.88	5.03	10.29	3.12	4.49	3.35	3.78	2.8	3.87
HERBA1003252	10.38	11.26	18.86	6.09	9.53	8.62	8.03	4.87	3.9
HERBA1003257	4.81	5.23	14.11	3.29	4.4	2.15	4.08	0.86	2.82
HERBA1003266	4.15	9.87	10.08	10.51	3.68	3.28	8.7	2.22	2.32
HERBA1003273	11.8	16.03	28.72	15.07	19.11	17.4	12.91	12.48	21.18
HERBA1003276	149.48	149.2	143.91	64.58	44.3	74.17	103.98	79.83	39.01
HERBA1003277	46.25	23.14	67.56	22.07	19.01	38.9	14.72	11.37	3.34
HERBA1003278	4.71	4.24	12.6	4.3	4.28	2.48	2.75	2.48	1.08
HERBA1003280	129.77	109.77	149.42	22.68	42.91	45.72	29.76	44.29	19.43
HERBA1003281	66.35	33.44	113.06	20.36	31.97	38.12	40.97	35.41	11.12
HERBA1003284	4.17	19.37	13.18	7.85	6.12	2.69	7.14	1.71	2.34
HERBA1003286	19.83	12.62	36.72	8.83	14.12	12.7	52.4	15.77	36.62
HERBA1003291	72.44	65.02	121.3	168.07	81.81	87.09	38.11	45.48	48.73
HERBA1003294	2.81	2.31	6.5	6.88	2.5	2.72	2.18	1.36	1.58
HERBA1003296	44.35	42.6	76.86	89.38	63.17	84.95	32.4	28.59	38.37
HERBA1003304	9.89	11.83	11.34	8.37	7.94	10.02	7.8	7.09	8.26
HERBA1003306	13.02	19.68	19.63	24.93	17.19	18.48	10.47	8.08	13.83
HERBA1003309	1.78	3.09	6.44	4.38	2.65	4.86	2.49	0.95	2.28
HERBA1003314	2.38	3.32	2.13	4.28	4.38	3.35	2.16	1.3	2.04
HERBA1003316	4.7	9	9.09	4.13	2.95	4.64	2.23	3.29	3.07
HERBA1003322	21.61	21.29	38.71	38.62	23.41	31.81	14.65	15.44	18.23
HERBA1003326	6.02	5.69	11.31	6.59	5.79	6.51	2.11	1.05	2.65
HERBA1003327	7.54	6.06	15.43	8.47	7.53	8.8	4.99	4.52	3.75
HERBA1003328	105.65	106.38	147.91	185.04	127.61	111.83	74.7	76.39	78.34
HERBA1003330	8.01	12.33	12.91	7.65	8.96	7.41	6.19	2.7	6.68
HERBA1003340	3.04	4.48	6.33	4.48	9.58	12.27	5.11	0.11	2.03
HERBA1003369	4.61	6.54	9.12	7.83	6.84	2.91	5.15	1.7	5.2
HERBA1003370	3.59	5.98	8.68	5.87	3.43	4.27	2.85	5.19	3.68
HERBA1003373	11.8	17.95	18.19	6.05	9.3	7.7	3.82	6.22	4.16
HERBA1003376	3.97	3.38	4.86	3.82	1.97	2.14	1.54	0.79	1.25
HERBA1003380	1.93	2.67	6.1	1.31	0.26	1.87	0.83	1.72	0.89
HERBA1003384	26.49	38.34	57.67	29.22	29.1	27.24	18.27	17.42	21.24
HERBA1003387	16.13	19.1	24.1	17.73	14.37	13.92	9.34	7.52	11.3
HERBA1003392	2.68	3.97	8.74	2.76	4.49	2.32	2.44	0.39	3.04
HERBA1003396	2.69	3.42	9.32	1.54	4.93	3.88	3.02	1.02	1.64
HERBA1003399	20.63	28.8	52.54	29.45	22.45	24.07	12.89	13.2	21.21
HERBA1003400	3.66	5.58	6.33	3.68	4.46	4.23	1.87	2.26	1.87
HERBA1003402	38.86	50.8	85.59	22.6	36.87	47.3	19.41	16.83	28.09
HERBA1003403	4.18	3.18	7.45	2.51	2.47	2.16	1.4	2.61	3.27
HERBA1003408	25.83	33.63	38.04	42.18	33.38	41.52	23.7	19.86	28.4
HERBA1003412	5.21	8.25	13.71	2.82	6	4.22	2.63	1.63	2.45
HERBA1003417	21.02	35.52	34.24	13.8	14.17	10.78	5.28	6.88	9.43

14 / 157

2466

【0838】

【表367】

2467

HENBA1003418	9.48	10.25	16.29	13.88	11.67	7.88	6.52	7.48	6.95
HENBA1003420	15.13	21.37	34.64	8.98	7.12	8.99	3.93	4.32	8.38
HENBA1003425	3.65	4.02	7.55	3.03	3.24	2.46	1.29	1.12	2.83
HENBA1003433	12.41	15.45	35.09	18.37	16.85	15.28	10.85	6.12	17.58
HENBA1003440	22.96	34.38	33.51	20.78	28.45	21.38	23.78	22.34	28.48
HENBA1003442	12.86	16.85	24.45	11.09	13.33	10.29	12.08	9.47	11.31
HENBA1003447	27.49	33.89	33.23	17.4	23.73	19.49	24	17.15	19.81
HENBA1003453	4.92	8.64	13.34	4.01	5.69	5.13	2.36	4.48	3.88
HENBA1003461	31.42	55.4	52.29	25.18	39.87	29.85	35.26	28.74	37.73
HENBA1003463	20.03	30.11	90.85	47.21	1.41	0.9	30.58	24.08	33.93
HENBA1003465	4.42	6.66	9.68	7.8	5.17	6.44	3.4	2.84	6.1
HENBA1003480	8.18	9.01	15.32	7.58	7.38	7.85	10.22	5.28	8.1
HENBA1003485	8.87	6.95	9.95	4.16	3.42	4.03	5.45	1.61	3.1
HENBA1003487	40.58	64.18	72.24	24.53	33.37	41.02	45.43	44.42	48.48
HENBA1003492	47.07	69.53	47.91	15.87	68.78	43.2	54.01	24.37	34.94
HENBA1003494	8.97	13.82	17.69	5.4	7.03	7.88	9.14	5.83	9.54
HENBA1003497	27.59	34.3	36.93	14.93	24.2	20.37	27.88	19.63	25.35
HENBA1003503	17.58	24.95	1.08	0.82	1.26	0.04	6.53	10.32	21.23
HENBA1003511	10.13	12.8	29.58	12.03	11.2	10.45	9.65	8.73	14.36
HENBA1003528	10.75	13.8	9.65	5.22	4.34	5.54	2.88	3.04	2.97
HENBA1003530	6.02	6.68	12.41	4.68	4.85	4.75	5.24	2.55	3.96
HENBA1003531	12.73	22.2	30.85	9.86	14.37	9.03	13.98	8.08	18.14
HENBA1003532	13.85	14.91	22.65	9.4	21.22	7.2	8.43	8.83	11.91
HENBA1003538	10.23	12.68	27.62	8.9	13.63	10.34	9.88	5.53	8.3
HENBA1003545	4.35	9.29	13.49	7.12	5.88	8.1	7.44	2.65	6.35
HENBA1003546	6.63	3.47	17.38	20.28	8.26	12.59	2.98	5.15	4.61
HENBA1003548	2.71	6.58	8.78	3.2	9.19	3.14	5.24	4.03	2.98
HENBA1003553	4	4.81	7.29	3.53	3.84	2.73	2.82	2.01	2.64
HENBA1003555	8.02	14.32	16.54	7.4	9.1	7.1	7.78	7.09	7.03
HENBA1003556	5	8.35	8.92	2.85	4.91	3.37	2.9	2.25	2.94
HENBA1003560	11.54	14.47	20.06	8.38	13.16	7.52	7.48	5.19	5.3
HENBA1003565	12.47	6.48	14.4	4.78	7.63	5.19	7.77	3.47	4.1
HENBA1003588	187.82	373.84	257.14	82.85	183.28	98.06	64.68	26.85	13.07
HENBA1003589	19.02	23.29	63.64	40.15	29.03	28.38	20	16.6	26.95
HENBA1003571	8.12	11.43	9.83	6.26	10.52	5.78	8.95	5.27	5.01
HENBA1003579	22.13	20.38	39.63	17.84	15.09	12.59	17.53	15.68	17.94
HENBA1003580	18.63	22.77	31.1	39.35	20	21.57	12.72	8.11	11.25
HENBA1003581	4.54	7.44	11.61	2.84	2.83	4.01	3.4	1.33	1.8
HENBA1003591	9.77	30.24	26.57	6.75	16.55	5.72	11.87	5.17	9.59
HENBA1003596	11.6	18.67	38.48	13.28	20.4	12.94	13.52	5.6	8.35
HENBA1003597	34.44	63.28	65.19	37.08	35.21	30.06	23.33	8.38	16.41
HENBA1003598	2.25	2.74	3.02	2.41	2.07	1.24	1.39	0.8	1.38
HENBA1003600	21.91	28.68	48.28	21.72	21.88	19.48	19.9	18.88	27.48
HENBA1003602	5.44	4.97	13.36	4.53	6.21	5.48	3.61	3.18	3.61
HENBA1003604	4.52	2.89	9.05	3.91	3.29	1.58	2.58	2.6	3.04
HENBA1003610	13.16	15.85	27.61	10.3	9.81	5.68	10.32	8.77	9.07
HENBA1003615	14.12	19.51	37.31	8.32	15.31	10.64	25.53	7.2	9.53
HENBA1003617	2.15	10.78	12.38	1.63	5.11	1.57	4.87	0.26	2.23
HENBA1003620	1.75	4.53	12.78	1.34	3.78	1.52	1.5	0.27	0.68
HENBA1003621	12.05	18.03	37.29	21.98	18.08	16.37	10.88	5.38	14.38
HENBA1003622	7.68	6.59	10	3.27	5.47	4.78	6.87	4.55	2.99
HENBA1003630	43.11	47.17	88.26	32.33	48.01	44.24	29.83	21.05	20.55
HENBA1003637	920.88	472.86	880.3	385.52	804.5	411.7	621.11	978.32	238.58
HENBA1003640	62.65	69.88	125.99	45.58	48.68	42	65.31	45.48	45.21
HENBA1003645	15.97	22.75	34.7	18.9	17.55	14.87	21.4	9.84	18.1
HENBA1003648	5.76	13.27	28.88	11.35	10.28	4.41	12.2	6.82	10.03
HENBA1003647	68.54	91.82	142.01	91.35	74.78	80.56	83.06	37.92	60.33
HENBA1003656	41.62	48.94	78.5	39.68	28.08	37.31	18.8	12.53	17.55
HENBA1003662	9.43	10.68	20.41	18.54	13.88	13.03	9.85	4.93	11.88
HENBA1003666	8.44	8.9	21.02	18.11	19.64	14.67	11.75	6.74	12.93
HENBA1003667	27	32.49	46.33	29.7	40.13	34.43	25.82	21.77	25.87
HENBA1003670	16.71	15.12	35.71	10.11	19.12	14.79	10.97	18.94	10.08
HENBA1003674	3.06	3.93	11.88	12.14	7.89	5.81	4.08	1.48	5.42
HENBA1003677	3.62	4.3	9.92	2.94	5.25	2.08	4.03	0.72	1.84

15 / 157

2468

2469

2470

HENBA1003779	15.99	25.43	34.72	23.29	23.03	14.07	32.67	15.26	20.68				
HENBA1003780	2.36	1.32	3.29	2.39	1.65	4.28	2	1.23	1.59				
HENBA1003784	27.44	32.04	37.17	24.58	15.64	21.48	12.03	8.81	13.69	*	**	-	-
HENBA1003790	28.38	32.22	54.74	28.94	29.04	32.33	34.38	26.69	39.53				
HENBA1003792	6.49	10.11	9.34	7.04	5.16	6.04	7.51	6.23	5.84				
HENBA1003792	8.66	9.3	12.43	12.4	14.68	13.17	8.17	5.02	7.69				
HENBA1003711	8.11	10.45	15.18	11.98	9.29	6.3	7.03	5.46	7.91				
HENBA1003714	8.44	11.03	12.83	2.97	13.48	9.98	7.88	7.19	9.67				
HENBA1003716	14.81	13.59	14.58	12.43	13.78	10.5	10.14	8.94	12.29	*		-	-
HENBA1003717	42.45	55.77	82.27	41.74	41.39	40.14	43.48	40.44	41.67				
HENBA1003720	6.21	6.28	13.24	9.93	12.72	5.69	6.86	6.06	6.12				
HENBA1003725	3.95	3.77	7.26	6.08	3.65	3.38	2.78	2.23	2.52				
HENBA1003726	4.28	12.28	8.88	10.99	2.88	4.38	6.4	1.69	3.67				
HENBA1003729	12.13	13.18	16.13	8.92	13.67	11.13	18.08	6.56	14.8				
HENBA1003732	30.17	53.31	40.17	33.85	38.01	30.07	19.05	11.75	17.73	*		-	-
HENBA1003733	3.45	4.52	4.35	2.89	6.7	2.94	3.18	1.76	2.64	*		-	-
HENBA1003742	9.01	9.99	10.98	9.58	10.71	5.69	9.3	5.8	8.68				
HENBA1003743	9.87	14.77	20.42	19.63	13.74	11.37	9.02	7.5	9.38				
HENBA1003750	14.27	20.31	26.23	15.6	15.1	13.01	12.22	13.09	13.08				
HENBA1003760	19.21	17.1	29.61	6.01	14.64	12.38	3.4	6.8	6.12	*		-	-
HENBA1003764	14.01	17.88	25.78	13.37	20.6	16.59	15.38	15.5	19.31				
HENBA1003769	39.51	54.43	53.68	30.85	34.08	32.37	42.08	31.8	48.8	*		-	-
HENBA1003773	5.18	9.28	8.37	4.89	7.62	4.92	2.95	1.8	4.58	*		-	-
HENBA1003783	2.33	2.73	7.18	5	7.98	9.31	5.24	2.04	3.68				
HENBA1003784	24.68	20.11	23.17	22.69	20.48	14.65	15.38	8.3	11.94	*		-	-
HENBA1003794	35.77	59.63	74.01	32.35	32.73	29.52	23.99	30.47	35.84				
HENBA1003799	28.51	43.08	44.43	24.39	25.48	22.11	20.76	21.74	28.31	*	*	-	-
HENBA1003803	6.8	7.68	12.5	2.71	3.98	6.01	1.91	2.94	2.14	*	*	-	-
HENBA1003804	12.17	15.5	24.58	5.01	10.25	5.91	7.51	11.06	8.61				
HENBA1003805	15.63	21.11	29.95	13.7	15.15	11.69	16.1	12.06	15.78				
HENBA1003807	12.24	15.49	18.72	15.37	14.25	10.11	18.36	12.3	19.08				
HENBA1003810	4.78	5.92	11.21	4.53	8.91	7.45	6.79	2.53	4.42				
HENBA1003827	18.51	13.97	18.38	12.97	15	8.61	18.16	7.58	9.11				
HENBA1003836	4.91	7.17	10.85	4.73	8.6	8.63	3.27	4.61	6.48				
HENBA1003838	49.74	66.87	78.01	40.91	58.53	48.34	31.6	28.8	30.51	*		-	-
HENBA1003842	23.09	38.55	55.11	24.97	28.81	27.21	27.17	12.95	30.84				
HENBA1003846	18.74	19.56	28.45	28.4	19.76	21.25	14.77	12.94	15.9				
HENBA1003856	31.17	48.34	48.82	28.74	30.87	25.87	49.83	42.07	42.91				
HENBA1003857	3.66	4.84	10.52	2.38	4.43	1.85	3.66	2.4	2.28				
HENBA1003864	24.35	38.18	29.59	24.03	32.07	18.61	15.17	4.79	6.89	*		-	-
HENBA1003866	15.55	24.7	24.92	21.6	24.17	13.68	22.15	14.95	14.18				
HENBA1003888	3.42	4.24	4.55	5.4	4.68	2.62	1.88	2.32	3.07	*		-	-
HENBA1003879	54.65	71.41	89.07	52.85	60.23	59.91	38.15	32.08	60.85				
HENBA1003880	26.75	40.67	50.47	32.31	30.67	29.71	28.74	17.94	31.69				
HENBA1003884	3.51	2.7	10.4	2.34	3.18	1.65	5.36	1.65	2.02				
HENBA1003885	15.83	16.07	23.65	9.67	12.61	9.91	13.22	10.04	13.72	*		-	-
HENBA1003887	8.92	10.53	15.63	3.65	12.51	2.64	7.43	4.94	3.2				
HENBA1003890	5.77	11.6	13.5	3.76	5.14	4.95	4.38	2.51	3.42	*		-	-
HENBA1003893	2.6	2.24	6.97	3.71	6.34	3.91	2.06	0.65	2.78				
HENBA1003896	13.18	13.74	38.07	17.77	20.7	14.83	10.45	11	13.75				
HENBA1003902	6.09	5.71	13.87	4.58	5.68	4.39	3.52	2.56	5.08				
HENBA1003904	41	55.7	67.32	43.67	30.73	35.35	31.08	28.57	32.22	*		-	-
HENBA1003908	27.42	42.25	49.6	27.37	22.92	19.91	28.26	21.38	25.22				
HENBA1003926	35.54	56.98	89.19	56.48	59.31	44	48.79	37.72	60.83				
HENBA1003937	14.37	14.68	20.38	10.43	17.34	12.15	14.47	8.13	16.59				
HENBA1003939	3.4	3.02	10.99	1.88	5.36	0.25	1.12	0.35	2.25				
HENBA1003940	6.76	11.65	10.45	4.59	10.58	5.65	9.9	2.16	3.94				
HENBA1003941	31.48	48.87	93.42	58.75	59.75	51.31	23.68	24.05	38.11				
HENBA1003942	3.19	4.4	9.44	4.64	4.07	4.72	1.95	2.11	3.24				
HENBA1003945	59.01	80.97	80.84	37.7	37.04	48.08	37.87	33.83	28.54	*	**	-	-
HENBA1003948	12.02	16.53	20.22	10.35	8.89	9.69	11.57	6.85	10.8				
HENBA1003950	27.53	47.63	43.7	25.63	28.11	14.15	20.64	11.87	12.95	*		-	-
HENBA1003963	10.33	12.12	25.01	8.15	9.43	8.53	13.65	9.93	7.98				
HENBA1003958	4.65	6.68	15.41	6.73	7.93	3.72	5.9	3.43	4.74				

16 / 157

2471

2472

MEMBA1003958	14.89	18.63	30.66	18.6	18.34	12.43	19.48	10.78	14.94				
MEMBA1003960	6.1	5.31	11.07	8.55	9.98	6.82	4.38	4.38	3.84				
MEMBA1003964	4.32	4.38	6.33	4.53	3.74	1.94	2.49	2.11	0.85				
MEMBA1003967	11.38	13.34	18.36	11.07	6.8	6.77	8.93	7.19	6.9				
MEMBA1003968	62.83	102.28	87.2	14.25	28.41	18.31	16.78	9.02	11.36				
MEMBA1003974	4.37	4.4	7.25	2.02	4.53	1.13	3.82	0.89	2.58				
MEMBA1003976	15.15	20.18	27.02	12.74	18.84	11.73	18.38	14.83	18				
MEMBA1003977	8.78	9.42	22.28	9.84	11.17	3.97	8.14	5.18	9.28				
MEMBA1003978	9.01	15.51	20.07	10.86	8.71	7.75	11.84	5.73	13.48				
MEMBA1003981	55.09	77.17	117.23	62.81	61.44	47.41	51.47	46.06	74.31				
MEMBA1003982	5.4	13.31	18.21	3.94	11.62	6.14	3.92	4.82	3.97				
MEMBA1003985	7.59	8.14	15.01	6.55	7.27	6.74	5.12	5.29	7.02				
MEMBA1003987	10.14	14.01	20.78	6.87	9.2	5.37	6.58	6.5	8.98				
MEMBA1003989	10.11	15.44	20.52	5.53	5.4	4.88	8.68	4.07	5.03				
MEMBA1004000	2.93	4.86	11.45	2.25	2.97	1.78	2.52	0.75	2.64				
MEMBA1004006	18.54	20.18	42.2	18.19	19.39	15.33	18.84	12.3	17.01				
MEMBA1004007	19.65	39.73	40.28	28.82	20.72	17.41	38.95	21.37	27.19				
MEMBA1004010	87.05	91.68	2.48	0.19	0.95	0.35	50.08	44.9	72.79				
MEMBA1004011	10.54	11.89	17.43	5.17	7.48	5.81	5.72	5.31	4.01				
MEMBA1004012	20.42	22.3	18.75	7.27	17.23	8.75	11.84	7.16	6.96				
MEMBA1004015	4.28	3.69	8.92	3.2	3.73	2.56	1.97	1.37	3.1				
MEMBA1004024	7.24	8.25	20.14	4.09	6.98	4.29	7.49	2.93	4.38				
MEMBA1004029	8.05	5.65	11.19	0.74	1.77	1.68	2.55	0.21	0.47				
MEMBA1004035	30.88	38.16	85.02	18.09	23.34	18.57	34.1	18.52	13.77				
MEMBA1004042	0.37	2.38	5.48	0.32	1.77	1.14	0.62	0	0.22				
MEMBA1004045	20.4	33.63	50.23	28.78	28.43	29.29	14.02	11.2	16.1				
MEMBA1004046	37.85	38.8	52.89	50.28	40.74	37.18	34.52	20.41	13.02				
MEMBA1004048	21.48	22.41	47.43	24.15	22.42	24.41	19.84	20.47	15.98				
MEMBA1004051	9.9	8.46	21.25	12.84	10.63	10.43	12.39	7.87	8.34				
MEMBA1004053	15.29	25.2	37.62	19.88	13.35	13.99	21.18	11.93	18.88				
MEMBA1004055	5.25	11.05	5.64	0.16	1.02	0	8.11	3.17	5.22				
MEMBA1004056	2.41	5.85	10.01	3.45	3.17	2.37	4.07	0.71	2.14				
MEMBA1004060	8.43	8.34	15.71	14.49	7.75	8	14.41	5.93	6.72				
MEMBA1004061	7.43	8.52	15.21	11.54	9.65	13.28	5.83	3.71	6.22				
MEMBA1004067	8.87	11.51	14.78	16.09	9.22	11.3	7.43	5.81	9.07				
MEMBA1004071	2.92	2.4	5.93	4.14	2.31	3.09	3.1	0.97	2.2				
MEMBA1004074	128.78	177.45	131.41	229.38	180.88	89.8	187.87	113.1	134.85				
MEMBA1004078	20.55	20.84	34.63	28.09	31.27	21.15	16.53	18.06	27.35				
MEMBA1004085	20.65	31.8	30.03	32.27	21.47	13.85	11.87	12.81	9.58				
MEMBA1004086	2.06	3.78	5.23	2.03	3.33	2.95	2.57	0.6	1.89				
MEMBA1004097	1.94	3.31	1.58	1.13	4.7	3.06	2.83	1.2	1.23				
MEMBA1004100	14.77	24.98	29.01	20.87	18.7	13.86	10.94	12.24	16.57				
MEMBA1004103	7.32	13.02	15.94	12.18	10.55	9.17	7.19	6.37	7.48				
MEMBA1004110	19.03	28.3	37.33	22.14	20.14	20.02	18.84	16.81	19.14				
MEMBA1004111	24.42	41.97	41.29	22.57	21.56	20.81	28.72	16.88	19.8				
MEMBA1004112	10.47	10.61	15.29	11.7	10.51	11.56	7.45	6.39	9.82				
MEMBA1004130	6.32	9.6	13.12	7.07	7.66	3.98	10.02	6.52	8.94				
MEMBA1004131	11.62	15.68	22.49	12.3	15.58	10.2	7.69	8.91	12.88				
MEMBA1004132	1.7	3.68	8.07	2.75	4.82	2.5	5.8	1.43	6.28				
MEMBA1004133	61.09	100.87	108.22	48.84	25.58	18.84	11.83	38.24	13.37				
MEMBA1004136	8.15	8.84	9.84	8.72	6.31	5.94	4.18	4.48	5.03				
MEMBA1004143	8.39	11.32	13.28	4.24	5.74	4.33	4.63	4.09	3.58				
MEMBA1004146	12.23	19.23	24.8	16.14	12.63	10.61	14.04	14.11	16.75				
MEMBA1004148	5.02	6.85	14.92	7.38	6.63	8.78	2.34	5.06	6.01				
MEMBA1004149	13.01	27.48	21.69	12.24	21.38	25.46	6.33	6.45	12.24				
MEMBA1004150	182.3	225.69	242.62	246.24	265.35	201.29	181.22	151.68	208.57				
MEMBA1004154	54.95	73.81	77.68	67.59	81.25	43.45	53.14	35.47	45.88				
MEMBA1004164	27.89	39.95	75.25	53.13	38	41.78	27.04	21.48	32.68				
MEMBA1004168	27.29	41.16	48.6	30.28	31.63	25.87	20.87	18.25	25.4				
MEMBA1004199	11.53	17.04	30.48	12.77	16.45	14.72	15.81	10.26	14.34				
MEMBA1004200	10.21	15.6	23.4	12.76	12.18	11.22	9.51	8.03	10.57				
MEMBA1004201	27.51	39.39	51.71	29.35	27.61	28.68	24.84	21.71	24.17				
MEMBA1004202	18.3	30.46	37.1	17.92	20.38	16.33	20.83	18.59	21.88				
MEMBA1004203	11.35	16.48	25.18	12.27	10.37	8.08	12.69	8.41	9.37				

17 / 157

2473

HENBA1004207	5.23	7	7.18	5.06	6.93	4.27	6.27	2.78	3.75				
HENBA1004210	4.96	5.53	8.99	5.1	8.24	7.88	8.78	4.25	8.82				
HENBA1004225	11.76	15.11	16.08	1.96	9.21	5.85	1.74	2.95	4.57	*	00	-	-
HENBA1004227	5.13	5	14.56	8.73	5.83	3.8	3.56	2.62	3.47				
HENBA1004235	55.87	87.38	100.88	50.82	83.74	84.88	58.93	44.28	81.25				
HENBA1004237	83.08	67.44	87.85	54.44	56.68	54.92	61.37	42.55	49.6	**		-	-
HENBA1004238	5.4	9.74	19.37	3.19	7.79	5.08	4.65	3.62	5.8				
HENBA1004241	58.53	84.19	86.84	45.42	81.25	30.71	41.3	31.18	35.09	*		-	-
HENBA1004242	95.96	84.64	132.52	129.29	124.32	88.47	88.28	40.26	48.03				
HENBA1004243	7.18	9.59	11.11	5.01	7.55	7.79	4.07	6.19	8.9				
HENBA1004246	3.92	5.45	6.92	3.64	1.73	7.81	2.23	1.7	8.01	*		-	-
HENBA1004247	38.08	39.85	23.48	13.27	21.43	17.87	8.15	9.3	7.87	*	00	-	-
HENBA1004248	80.7	102.92	88.75	40.08	49.28	54.01	55.12	49.8	49.38				
HENBA1004250	5.48	7.1	17.92	5.82	8.67	3.92	3.52	3.06	5.68				
HENBA1004252	9.76	13.49	18.16	5.3	8	5.39	8.18	4.43	8.93	*		-	-
HENBA1004260	3.71	7.97	10.5	2.84	9.75	1.75	6.82	1.65	1.83				
HENBA1004264	4.86	4.92	5.04	3.34	5.85	4	4.08	2.37	1.97	*		-	-
HENBA1004272	14.31	28.05	30.38	18.35	16.25	13.91	15.22	15.98	20.33				
HENBA1004272	7.23	8.73	9.64	3.95	3.4	9.74	3.02	1.81	2.67	**	00	-	-
HENBA1004274	3.8	4.23	13.32	4.59	3.82	2.88	5.15	2.11	5.65				
HENBA1004276	6.92	5.28	11.01	3.44	3.27	3.1	4.82	3.75	4.58				
HENBA1004276	53.88	104.51	88.12	40.54	57.2	47.57	40.18	23.52	29.44	*		-	-
HENBA1004278	3.9	4.4	11.72	1.47	4.01	1.18	2.34	1.51	2.01				
HENBA1004284	15.6	17.68	32.1	11.85	15.45	12.28	19.44	11.68	15.92				
HENBA1004285	1.1	4.62	5.35	1.91	3.11	0	3.7	0.1	2.34				
HENBA1004289	4.33	10.31	14.89	4.93	9.21	6.85	5.43	4.43	9.27				
HENBA1004293	14.22	23.33	31.53	18.23	14.77	12.78	18.77	15.46	15.6				
HENBA1004295	5.99	9.65	12.39	6.69	4.97	4.93	6.51	6.63	6.32				
HENBA1004302	27.72	69.78	67.78	35.38	30.81	31.88	31.74	25.34	24.05				
HENBA1004308	22.59	41.69	30.11	30.27	34.77	20.17	21.64	11.35	18.3				
HENBA1004312	8.29	12	23.23	5.75	9.79	3.74	7.73	4.58	8.86				
HENBA1004314	17.03	23.79	39.64	14.71	20.85	13.02	20.7	18.59	24.72				
HENBA1004321	2.18	4.08	8.11	3.21	4	1.12	2.54	0	3.38				
HENBA1004323	4.43	3.01	0.88	0.1	10.07	1.6	2.71	1.11	0.96				
HENBA1004327	3.2	6.57	7.13	3.58	3.76	2.03	3.61	2.13	4.27				
HENBA1004329	47.09	53.32	108.52	58.12	44.85	40.88	71.49	47.58	48.41				
HENBA1004330	7.28	14.18	16.85	8.08	8.07	7.74	11.04	8.62	5.53				
HENBA1004334	6.28	6.88	13.5	5.07	8.47	3.23	5.33	2.42	4.72				
HENBA1004335	2.77	3.49	11.85	1.87	3.84	0.75	1.36	0.2	2				
HENBA1004341	4.79	7.07	13.59	3.55	3.83	2.11	5.23	1.8	5.75				
HENBA1004344	5.88	15.53	22.86	10.54	9.43	7.82	16.14	4.47	8.61				
HENBA1004347	8.48	17.07	21.37	11.89	14.38	7.58	5.28	6.18	12.85				
HENBA1004349	19.03	22.2	39.78	17.07	23.84	11.42	16.44	11.72	16.17				
HENBA1004352	30.35	31.05	54.47	23.2	27.22	18.91	26.94	20.35	19.47				
HENBA1004353	65.35	90.37	111.41	40.12	57.25	44.69	82.88	49.41	51.19	*		-	-
HENBA1004354	31.04	48.91	55.45	29.8	25.92	23.18	26.7	16	25	*	0	-	-
HENBA1004356	38.8	69.45	87.9	42.77	38.63	30.42	48.78	31.27	34.28				
HENBA1004360	21.37	39.1	80.88	24.3	25.32	19.97	35.28	18.19	17.1				
HENBA1004366	3.38	19.8	13.89	9.13	8.25	3.68	5.68	3.78	6.29				
HENBA1004372	26.71	35.24	61.82	27.78	27.86	28.9	21.52	16.32	31.11				
HENBA1004377	57.43	64.54	125.12	52.77	52.38	39.76	63.75	39.09	45.41				
HENBA1004389	28.8	23.69	66.26	35.4	23.53	18.76	21.22	13.58	17.27				
HENBA1004391	15.02	20.13	38.52	17.1	15.91	14.1	20.59	16.16	16.69				
HENBA1004393	8.23	8.43	14.77	10.01	2.64	5.81	6.08	4.78	4.24				
HENBA1004394	6.59	9.68	22.73	7.13	7.38	4.87	8.85	2.4	3.32				
HENBA1004396	6.88	7.79	21.04	5.3	10.85	6.63	4.48	2.98	4.78				
HENBA1004401	25.42	37.64	61.19	35.6	34.11	30.34	50.39	18.8	33.72				
HENBA1004405	12.61	20.04	37.41	22.75	14.99	14.73	16.19	9.3	19.54				
HENBA1004408	33.43	47.3	70.7	34.73	38.57	32.16	33.47	25.45	21.56				
HENBA1004414	11.75	12.41	34.08	20.66	20.73	15.48	9.98	9.26	12.21				
HENBA1004429	11.38	11.1	25.62	16.77	12.96	11.07	10.66	8.23	8.04				
HENBA1004433	4.53	4.23	9.92	6.93	6.74	8.38	4.52	0.98	3.02				
HENBA1004440	12.02	23.81	31.06	19.87	29.44	19.7	15.12	10.66	15.31				
HENBA1004444	18.31	24.79	48.64	21.84	22.35	18.04	25.64	13.85	21.08				

18 / 157

2474

[0842]

[表371]

2475

2476

HEIBA1004440	1.00	3.03	4.89	2.99	6.41	1.98	3.27	0.17	1.81
HEIBA1004441	47.91	49.66	68.71	31.81	23.94	40.91	29.4	25.22	31.73
HEIBA1004452	7.96	9.5	12.83	8.01	5.07	6.5	4.85	4.18	4.59
HEIBA1004454	21.61	21.48	36.91	22.76	21.2	22.64	20.86	13.58	20.72
HEIBA1004460	3.23	5	9.05	6.23	2.51	4.45	3.98	1.85	1.76
HEIBA1004481	9.82	11.87	18.26	13.48	12.48	12.25	11.75	5.13	11.97
HEIBA1004488	5.97	8.24	11.49	13.09	9.63	6.4	4.93	3.14	5.27
HEIBA1004479	7.22	6.26	11.32	4.91	8.01	9.21	5.84	3.77	4.35
HEIBA1004482	5.37	8.99	11.74	7.23	3.06	5.24	4.77	2.99	3.49
HEIBA1004481	7.67	10.41	13.9	11.84	11.36	7.74	9.41	6.84	7.89
HEIBA1004489	7.91	12.25	21.91	14.4	10.84	11.41	12.99	9.9	12.68
HEIBA1004502	28.69	31.67	44.63	24.39	21.51	21.75	15.45	9.07	14.65
HEIBA1004505	9.98	9.29	16.72	11.78	9.95	10.23	12.78	4.89	8.35
HEIBA1004506	10.31	10.67	16.89	10.45	12.56	11.51	11.7	4.69	11.13
HEIBA1004507	21.1	25.48	30.3	21.45	19.73	17.8	29.74	17.99	32.44
HEIBA1004509	2.41	4.16	3.19	4.33	6.97	5.13	3.86	1.09	3.24
HEIBA1004523	18.14	17.44	35.11	23.55	14.39	16.02	17.09	11.42	16.4
HEIBA1004528	10.51	13.24	19.76	9.67	18.03	13.29	8.5	10.38	8.52
HEIBA1004534	16.22	16.88	30.37	15.84	14.03	13.54	14.2	7.84	17.71
HEIBA1004538	6.18	6.2	10.88	7.87	5.18	6.58	4.93	4.17	3.26
HEIBA1004539	6.62	8.85	16.21	10.28	7.48	8.38	4.46	3.25	4.87
HEIBA1004542	7.28	10.06	13.66	7.78	7.83	8.12	4.08	4.75	6.99
HEIBA1004552	3.28	4.65	8.81	4.78	5	6.2	4.33	0.53	3.84
HEIBA1004584	6.18	11.95	16.82	9.67	13.98	10.14	9	4.19	10.19
HEIBA1004589	12.38	17.94	21.39	13.09	11.7	13.96	9.84	8.22	10.93
HEIBA1004600	4.66	8.98	18.88	4.85	6.89	4.95	2.88	5.27	5.35
HEIBA1004564	9.73	17.48	27.38	18.06	14.28	12.12	13.33	10.15	10.66
HEIBA1004588	3.56	3.98	12.25	2.38	4.04	2.92	3.05	2.38	2.63
HEIBA1004573	4.48	4.38	12.03	4.58	4.29	4.64	2.63	3.89	4.43
HEIBA1004576	18.09	17.82	32.93	24.31	34.17	25.83	17	11.99	13.66
HEIBA1004577	3.64	7.83	16.24	7.08	11.02	14.75	9.75	3.87	4.69
HEIBA1004586	4.65	5.81	14.36	5.67	7.68	6.89	6.37	2	4.74
HEIBA1004594	284.44	249.12	425.82	413.26	281.05	315.49	138.33	269.7	335.01
HEIBA1004604	11.58	13.47	19.78	11.02	11.29	12.05	9.52	8.82	11.26
HEIBA1004607	44.74	50.81	68.68	61.55	58.7	51.72	48.7	32.51	58.83
HEIBA1004610	6.05	7.49	14.69	9.25	6.9	7.29	6.72	4.7	5.16
HEIBA1004617	12.1	9.56	20.48	8.02	10.18	8.41	9.52	5.32	9.88
HEIBA1004622	5.18	6.28	10.24	4.13	6.32	4.27	7.06	2.49	4.07
HEIBA1004626	5.67	7.39	15.1	4.2	9.2	4.95	4.59	5.69	4.1
HEIBA1004629	13.02	21.22	41.46	18.54	21.07	18.99	20.74	13.12	14.82
HEIBA1004631	73.82	82.05	86.06	51.43	28.56	15.5	10.72	28.9	42.8
HEIBA1004632	24.68	42.63	52.35	37.34	39.22	29.29	28.86	22.05	34.67
HEIBA1004633	38.98	38.11	53.52	17.84	40.68	39.18	15.84	33.56	35.48
HEIBA1004636	18.64	14.75	24.41	11.5	12.22	12.63	6.1	4.97	7.02
HEIBA1004637	19.33	12.11	26.18	6.02	8.26	11.39	12.44	5.03	12.85
HEIBA1004638	17.07	21.26	52.09	19.97	27.34	24.5	24.76	14.24	22.8
HEIBA1004645	41.39	65.54	74.75	36.73	58.12	57.87	35.72	21.96	40.06
HEIBA1004656	33.89	68.4	60.19	35.73	48.51	37.77	45.77	32.71	37.95
HEIBA1004657	8.65	8.21	14.8	8.65	5.01	4.76	2.7	2.34	4.48
HEIBA1004656	18.27	17.29	28.67	14.15	16.18	11.73	13.41	11.88	17.32
HEIBA1004689	9.31	10.17	13.58	7.96	9.32	5.8	4.8	3.37	4.61
HEIBA1004670	0.26	8.23	12.51	5.31	4.65	7.13	5.07	1.93	2.73
HEIBA1004672	4.49	6.42	13.8	6.32	5.91	7.14	7.17	1.42	6.19
HEIBA1004688	18.08	37.4	36.02	13.85	14.84	13.32	10.59	5.45	12.38
HEIBA1004690	8.71	10.59	14.84	2.74	5.28	4.9	4.08	1.25	4.02
HEIBA1004693	64.63	61.62	109.88	81.26	88.47	81.99	74.06	35.68	58.9
HEIBA1004697	57.99	90.64	174.23	129.1	114.25	71.95	45.95	87.21	109.42
HEIBA1004702	17.82	20.4	33.62	24.43	14.82	16.84	15.1	13.79	12.53
HEIBA1004704	4.12	4.28	3.85	3.01	2.64	2.5	2.8	1.84	1.97
HEIBA1004705	8.63	8.58	10.66	4.87	5.69	4.53	7.25	3.35	5.23
HEIBA1004706	4.44	4.15	10.31	3.38	3.58	2.66	3.1	1.58	2.81
HEIBA1004709	17.24	19.08	37.38	23.48	18.59	23.59	17.98	10.88	14.19
HEIBA1004711	162.08	157.28	293.47	260.03	204.85	251.04	138.17	104.82	95.04
HEIBA1004723	8.81	12.37	23.58	15.07	18	14.65	13.6	6.03	7.35

2477

HENBA1004725	8.16	9.51	14.17	11.97	14.28	12.48	9.22	7.11	10.71										
HENBA1004730	9.5	30.77	15.01	11.4	24.39	6.98	5.89	6.95	7.79										
HENBA1004733	4.35	1.84	9.11	2.44	2.1	1.31	3.16	0.68	3.23										
HENBA1004734	6.35	4.58	8.64	3.47	5.46	3.65	7.83	2.85	5.88										
HENBA1004735	16.7	24.9	39.79	8.08	17.83	7.89	15.58	8.45	13.06										
HENBA1004748	7.15	9.02	16.19	2.15	7.19	3.48	6.72	4.3	8.2										
HENBA1004751	24.38	23.03	33.63	4.23	9.92	4.61	5.83	2.01	3.8	**	**	-	-						
HENBA1004752	2.98	5.64	15.55	3.81	7.93	4.03	5.98	2.05	2.94										
HENBA1004753	114.05	106.99	178.18	68.94	72.11	86.48	30.58	32.81	51.61										
HENBA1004755	63.77	72.42	98.41	28.89	47.7	46.78	41.85	35	54.38	*	*	-	-						
HENBA1004756	14.23	13.71	21.69	11.51	11.62	10.29	10.63	11.28	11.46										
HENBA1004758	8	8.08	13.15	9.58	10.37	12.14	11.47	5.24	11.51										
HENBA1004763	28.52	33.28	48.87	20.49	28.34	19.64	31.58	21.08	23.78										
HENBA1004765	5.65	6.08	11.45	2.37	10.09	17.87	15.26	61.35	5.43										
HENBA1004768	3.64	9.18	17.67	4.3	5.33	4.44	3.13	1.14	2.86										
HENBA1004770	3.52	3.97	15.56	4.59	3.64	3.31	3.4	1.61	4.49										
HENBA1004771	9.14	8.57	13.33	7.83	4.79	6.17	4.13	3.12	6.19	*		-	-						
HENBA1004775	4.31	4.72	7.01	2.54	3.11	3.88	4.57	1.14	2.89										
HENBA1004776	24.15	24.27	34.87	11	13.72	16.67	11.79	13.87	5.14	*	*	-	-						
HENBA1004778	7.73	3.88	15.13	3.29	12.02	8.24	8.23	3.84	2.81										
HENBA1004784	24.44	38.5	43.33	28.54	19.27	15.1	33.56	28.4	19.49										
HENBA1004785	13.67	18.37	27.67	13.73	15.71	13.71	15.09	7.19	15.39										
HENBA1004789	4.14	8.07	13.1	2.68	4.24	5.12	10.75	3.53	4.53										
HENBA1004795	7.33	8.35	20.45	14.48	8.52	7.53	6.22	3.52	6.52										
HENBA1004797	7.1	4.75	10.25	8.19	4.85	6.24	4.01	3.13	5.6										
HENBA1004803	8.6	9.94	11.35	7.93	6.31	7.52	6.11	2.35	5.54	*	*	-	-						
HENBA1004806	3.19	2.79	8.69	6.88	5.28	4.11	3.33	0.92	0.9										
HENBA1004807	4.03	3.35	11.64	5.63	6	7.16	2.01	2.6	2.3										
HENBA1004818	15.48	10.87	19.39	6.27	7.54	6.26	5.63	6.15	5.23	*	*	-	-						
HENBA1004820	7.87	10.12	20.63	12.23	11.52	10.1	9.12	3.16	8.37										
HENBA1004833	4.21	6.73	18.25	5.9	8.04	9.77	5.85	1.88	3.49										
HENBA1004847	3.02	6.46	11.24	4.65	7.78	8.68	8.67	2.6	5.11										
HENBA1004850	4.4	5.74	10.78	6.32	4.58	7.77	3.54	4.67	2.65										
HENBA1004863	17.31	15.88	32.11	21.23	15.33	16.21	14.88	9.41	17.84										
HENBA1004864	17.45	20.24	32.03	15.38	13.18	16.06	19.67	7.45	13.87										
HENBA1004865	113.3	61.73	148.7	148.79	68.67	115.94	64.71	39.42	63.09										
HENBA1004880	8.83	11.17	12.28	9.73	8.92	10.05	5.47	2.68	3.9	**		-	-						
HENBA1004882	141.61	124.12	179	131.9	123.22	118.49	123.09	86.3	152.75										
HENBA1004885	7.94	7.93	12.26	7.41	11.08	9.32	5.37	3.13	5.59	*		-	-						
HENBA1004889	35.77	44.93	60.57	61.03	25.98	30.7	17.41	45.83	38.11										
HENBA1004900	52.96	49.82	61.95	21.61	27.4	15.4	7.06	13.28	12.57	**	**	-	-						
HENBA1004909	14.18	13.16	28.88	16.89	11.25	13.43	12.38	8.23	13.02										
HENBA1004918	6.03	4.11	8.13	4.43	3.48	3.67	3.79	1.63	1.47										
HENBA1004923	9.64	9.9	14.09	8.35	13.9	10.52	5.58	4.16	7.28	*		-	-						
HENBA1004929	3.15	3.03	7.44	3.03	3.05	3.22	2.76	0.8	3.27										
HENBA1004930	4.7	7.7	9.9	4.33	7.84	6.01	5.83	1.88	7.33										
HENBA1004933	48.8	50.84	85.65	52.38	60.44	49.38	52.85	24.63	54.95										
HENBA1004934	3.85	4.55	8.24	5.31	3.47	5.34	6.35	6.72	5.35										
HENBA1004937	12.55	13.42	25.29	16.02	10.93	15.03	11.34	9.36	13.18										
HENBA1004943	6.33	9.2	8.01	2.75	3.28	2.19	4.11	2.61	3.13	**	**	-	-						
HENBA1004944	19.58	18.31	31.82	17.17	17.7	18.72	15.82	10.66	19.2										
HENBA1004946	12.19	9.84	29.87	11.5	9.34	11.54	7.67	6.35	7.52										
HENBA1004952	5.62	5.6	7.61	2.28	4.07	4.82	1.95	1.52	2.27	**		-	-						
HENBA1004954	95.41	161.15	81.32	125.56	130.49	124.27	79.29	38.2	131.05										
HENBA1004958	3.24	5.15	10.1	3.76	7.59	6.47	4.82	2.28	4.38										
HENBA1004960	18.1	21	43	23.94	15.38	19.28	12.89	14.78	18.5										
HENBA1004971	17.31	28.71	28.24	5.08	6.02	5.94	4.63	4.76	6.7	**	**	-	-						
HENBA1004972	122.03	238.04	172.22	44.53	137.38	72.83	61.38	80.13	88.54	*		-	-						
HENBA1004973	3.44	2.83	7.72	3.58	1.82	6.45	2.6	2.08	3.24										
HENBA1004977	14	10.09	24.08	10.99	9.73	9.85	8.91	5.11	5.98										
HENBA1004978	11.81	9.75	18.48	9.68	8.28	8.27	8.68	4.77	6.85										
HENBA1004980	5.76	8.63	15.4	3.58	6.32	6.19	5.45	3.4	4.05										
HENBA1004982	8.24	8	15.9	4.88	8.72	7.61	5.71	2.32	4.42										
HENBA1004983	27.66	31.4	50.17	31.75	22.28	27.83	28.75	25.03	37.28										

20 / 157

2478

2480

MEMBA10050095	13.04	8.58	10.13	7.12	8.06	6.92	2.73	3.68	5.31
MEMBA10050094	47.77	5.38	6.34	4.11	3.26	3.94	3.94	3.84	5.52
MEMBA10050093	5.09	40.78	98.88	67.53	69.03	84.32	23.5	38.71	21.48
MEMBA10050092	1057.8	723.68	1508	888.08	1348.5	1172.9	1266.5	763.02	961.92
MEMBA10050091	4.91	4.1	3.37	12.38	2.46	5.05	1.17	5.8	3.17
MEMBA10050090	20.78	39.68	34.72	6.87	27.86	14.2	11.45	15.23	4.87
MEMBA10050089	70.70	80.68	114.47	109.76	103.35	90.04	59.53	40.68	63.52
MEMBA10050088	16.47	18.85	35.14	26.8	19.92	22.3	9.64	15.48	14.58
MEMBA10050087	8.48	8.45	12.94	4.31	7.1	4.1	3.89	2.47	4.91
MEMBA10050086	30.97	34.92	39.83	34.24	27.68	28.36	12.03	8.11	16.92
MEMBA10050085	8.28	10.14	13.81	4.91	5.09	5.36	2.77	2.57	2.31
MEMBA10050084	3.68	2.17	10.81	2.45	2.33	1.31	2.32	1.28	0.58
MEMBA10050083	14.16	12.91	14.13	4.05	3.84	22.08	6.52	3.12	5.74
MEMBA10050082	10.67	14.99	36.8	11.75	15.95	10.56	12.8	8.39	12.44
MEMBA10050081	4.44	5.11	10.87	2.4	4.78	3.1	2.95	0.22	2.1
MEMBA10050080	62.72	54.64	132.49	164.51	146.19	182.23	48.46	33.29	55.3
MEMBA10050079	3.89	4.62	7.08	4.3	4.86	3.7	4.12	2.52	4.18
MEMBA10050078	9.47	10.58	12.93	7.13	9.41	6.93	4.44	6.27	3.76
MEMBA10050077	31.02	35.09	45.38	24.48	22.17	19.63	32.37	21.21	24.67
MEMBA10050076	6.87	5.28	12.68	2.37	3.44	3.16	3.82	1.83	7.43
MEMBA10050075	14.72	21.03	25.93	7.18	18.88	15.76	10.99	5.75	6.49
MEMBA10050074	4.48	4.67	12.93	1.85	3.81	2.49	3.1	1.87	2.53
MEMBA10050073	4.71	9.88	14.88	4.88	6.06	3.51	4.1	0.92	1.92
MEMBA10050072	25.03	17.55	31.81	18.25	14.13	11.76	5.18	3.62	11.92
MEMBA10050071	5.89	3.91	4.85	4.65	2.95	3.4	2.39	2.2	2.94
MEMBA10050070	39.68	41.35	60.66	28.48	23.15	20.02	33.28	32.78	29.58
MEMBA10050069	40.1	40.38	55.64	28.82	24.01	24.18	41.74	30.86	33.16
MEMBA10050068	113.21	107.33	158.04	92.91	88.12	78.09	108.26	47.64	78.38
MEMBA10050067	10.25	16.19	27.19	10.38	11.81	8.06	14	8.52	9.01
MEMBA10050066	17.82	23.43	41.02	14.34	24.18	22.06	27.83	16.74	26.23
MEMBA10050065	5.58	8.93	16.2	4.63	7.32	4.29	4.52	1.48	2.68
MEMBA10050064	6.53	2.71	8.45	5.36	4.35	4.23	3.05	2.29	4.35
MEMBA10050063	378.91	317.85	317.18	22.18	181	206.94	102.53	216.48	124.98
MEMBA10050062	8.62	9.2	10.54	4.39	6.2	1.79	3.69	2.97	2.92
MEMBA10050061	7.43	8	10.73	4.97	8.07	2.08	4.16	3.29	1.57
MEMBA10050060	7.46	10.34	13.32	4.47	4.62	4.49	4.3	1.57	2.99
MEMBA10050059	22.87	28.13	43.59	20.68	28.1	16.31	27.6	18.67	18.78
MEMBA10050058	8.28	12.01	22.97	7.71	8.71	6.32	6.16	2.19	5.11
MEMBA10050057	21.95	45.46	54.34	18.83	21.77	13.25	11.91	3.42	10.22
MEMBA10050056	38.23	25.81	53.48	38.65	31.34	27.12	18.26	15.51	22.99
MEMBA10050055	5.69	7.58	9.58	3.99	3.6	4.2	6.81	2.2	7.06
MEMBA10050054	16.48	12.78	28.18	11.48	14.06	9.97	14.37	10.45	14.78
MEMBA10050053	23.26	6.89	27.27	4.57	14.1	5.14	8.86	8.11	3.99
MEMBA10050052	40.02	33.68	64.57	31.22	55.15	17.78	33.33	34.4	11.24
MEMBA10050051	8.36	5.89	13.54	4.44	2.76	4.32	3.73	1.82	7.18
MEMBA10050050	6.79	9.67	18.26	5.01	5.11	5.18	4.75	0.62	2.59
MEMBA10050049	211.84	132.19	314.59	37.27	92.18	87.58	104.99	40.98	12.2
MEMBA10050048	5.02	3.23	8	4.94	3.68	4.85	3.42	2.21	2.64
MEMBA10050047	4.01	4.87	12.04	12.05	4.27	6.06	2.73	1.86	3.31
MEMBA10050046	7.18	4.02	12.24	4.12	3.03	3.52	2.77	1.9	3.08
MEMBA10050045	4.98	6.13	14.8	5.66	6.84	4.51	2.63	3.68	2.47
MEMBA10050044	144.3	160.28	353.25	249	171.88	181.26	165.78	91.77	167.68
MEMBA10050043	11.91	8.88	32.76	5.64	10.3	10.41	18.87	11.34	4.4
MEMBA10050042	12.18	18.12	37.6	17.48	16.35	13.11	14.26	8.12	11.62
MEMBA10050041	42.25	46.46	69.97	65.43	48.12	53.85	34.92	14.48	29.09
MEMBA10050040	146.08	122.53	200.33	113.76	128.98	123.13	108.12	96.92	123.47
MEMBA10050039	50.95	62.4	71.8	62.62	44.87	45.43	64.52	35.24	62.21
MEMBA10050038	36.19	33.62	70.76	88.7	68.37	56.28	25.84	20.1	22.79
MEMBA10050037	11.95	9.22	34.15	19.57	13.03	24.04	13.57	18.82	7.5
MEMBA10050036	12.82	8.96	24.18	11.91	18.13	12.53	5.65	16.37	7.9
MEMBA10050035	7.64	4	12.18	6.44	7.08	7.49	6.65	3.04	4.73
MEMBA10050034	3.18	4.74	9.81	5.51	5.89	5.14	6.35	2.68	2.92
MEMBA10050033	4.23	5.99	12.03	5.04	5.33	5.38	6.54	0.63	2.54
MEMBA10050032	11.53	8.17	13.7	12.69	6.17	11	10.14	6.82	9.41

(1242)

特開2002-191363

2481

2482

HENBA10053113	33.27	32.37	38.89	32.29	19.06	24.67	17.8	18.01	18.82	**	-
HENBA10053114	24.58	33.06	42.65	21.87	11.43	15.5	9.82	9.33	8.85	*	-
HENBA10053115	12.58	12.96	19.94	17.38	10.31	13.38	9.84	8.48	5.8	*	-
HENBA10053117	4.9	4.68	7.03	6.55	5.75	6.41	3.83	2.01	4.56		
HENBA10053118	2.91	3.45	4.86	4.07	3.07	4.69	2.08	0.38	2.27		
HENBA10053224	11.54	12.48	27.19	15.78	17.61	18.74	11.73	1.86	14.23		
HENBA10053311	43.61	41.81	50.5	40.68	48.26	41.6	45.76	21.8	43.48		
HENBA10053317	20.3	20.68	34.18	18.47	13.27	15.91	13.2	7.95	13.18	*	-
HENBA10053338	37.76	51.08	62.83	38.88	14.3	16.63	9.18	11.59	18.35	**	-
HENBA1005344	58.27	53.4	127.13	158.88	86.48	122.88	43.73	24.53	38.87		
HENBA1005363	281.67	197.22	329.41	381.89	358.33	324.87	354.08	232.83	284.3		
HENBA1005369	3.54	4.47	8.65	4.81	7.38	8.25	4.29	2.88	3.92		
HENBA1005382	14.29	12.72	20.99	11.94	11.74	13.92	13.73	6.82	10.98		
HENBA1005384	3.08	8.57	11.19	9.32	9.41	5.54	6.78	3.23	3.82		
HENBA1005387	8.52	14.38	9.74	11.82	16.31	12.41	12.24	9.27	3.86		
HENBA1005372	209.34	404.69	430.49	84.13	123.89	78.11	25.56	68.19	97.12	**	**
HENBA1005374	35.56	51.63	59.8	44.44	39.19	38.53	38.51	36.03	34.64		
HENBA1005379	33.5	27	43.36	22.13	30.25	17.05	21.18	29.05	27.22		
HENBA1005382	15.49	19.68	32.6	16.35	15.89	24.51	20.78	15.49	21.41		
HENBA1005384	6.38	8.47	15.31	11.14	12.78	12.40	8.42	6.41	12.29		
HENBA1005386	7.43	7.3	12.47	6.27	5.44	7.51	3.39	1.08	4.63	*	-
HENBA1005389	18.61	20.29	18.71	13.6	21.69	22.09	19.9	13.97	9.2		
HENBA1005394	12.89	8.67	16.27	12.3	14.5	11.32	10.11	3.58	3.69		
HENBA1005403	4.34	4.47	10.77	7.65	8.37	7.4	6.03	4.21	5.27		
HENBA1005408	5.14	5.28	8.12	4.84	5.38	6.7	5.4	4.72	1.33		
HENBA1005410	147.8	188.98	244.61	96.9	175.54	169.88	68.34	83.45	81.73	*	-
HENBA1005411	3.94	4.86	10.62	4.18	2.45	4.02	1.07	3.25	2.19		
HENBA1005423	7.19	6.73	19.41	6.38	7.84	8.56	6.14	3.21	5.83		
HENBA1005426	95.38	92.68	185.58	213.51	144.79	175.18	76.93	45.72	95.01		
HENBA1005427	3.18	3.68	8.55	2.9	4.08	4.42	2.98	1.86	3.86		
HENBA1005430	39.92	60.77	66.51	41.28	49.27	33.43	51.55	28.88	24.29		
HENBA1005438	85.73	96.52	156.38	59.22	45.61	57.38	12.78	14.31	32.68	*	-
HENBA1005443	15.2	22.18	31.05	12.38	21.42	15.9	15.18	10.34	15.89		
HENBA1005447	7.4	8.91	13.83	6.04	8.83	6.51	4.71	3.18	4.95	*	-
HENBA1005449	188.67	185.78	227.88	315.2	205.18	189.94	184.99	92.08	168.89		
HENBA1005452	87.1	62.51	97.54	153.23	76.5	108.72	73.14	41.42	77.68		
HENBA1005454	4.07	5.62	12.12	2.12	8.94	2.42	3.65	2.55	2.8		
HENBA1005488	3.12	5.06	12.85	2.9	5.78	4.72	3.8	1.69	1.54		
HENBA1005489	4.08	5.82	14.74	4.56	8.66	7.3	4.34	3.61	3.19		
HENBA1005472	48.84	68.93	98.37	43.92	45.46	42.18	48.44	44.29	63.31		
HENBA1005474	3.81	5.4	6.23	3.61	6.81	3.03	3.23	2.83	4.47		
HENBA1005475	18.35	13.18	28.74	13.89	11.52	8.61	14	8.88	16.92		
HENBA1005489	11.24	6.84	19.41	11.32	8.2	10.15	7.69	4.76	6.72		
HENBA1005497	3.55	3.84	13.2	3.87	5.02	4.47	2.68	1.64	6.08		
HENBA1005500	9.05	12.27	15.75	5.87	8.54	10.09	5.5	7.58	6.57	*	-
HENBA1005506	68.72	77.27	99.37	89.95	83.48	88.85	48.45	35.29	43.04	*	-
HENBA1005508	78.62	18.74	23.57	15.52	16.57	19.17	24.35	7.4	7.32		
HENBA1005511	186.43	145.35	110.4	80.22	48.21	48.23	5.8	10.19	8.56	**	**
HENBA1005513	13.22	13.61	12.43	5.61	8.6	9.42	6.35	6.5	6.97	*	**
HENBA1005517	8.83	13.6	21.17	12.19	8.09	9.65	12.08	10.72	8.14	*	*
HENBA1005518	6.15	6.12	10.29	3.31	2.65	3.34	2.82	2.51	2.48		
HENBA1005520	4.72	5.27	15.68	1.8	8.98	5.07	11.44	4.1	6.1		
HENBA1005522	7.43	8.22	15.1	2.89	4.51	4.58	4.2	1.89	5.13		
HENBA1005526	12.95	11.98	23.79	8.97	14.67	6.23	7.41	3.51	8.41		
HENBA1005528	28.88	31.78	47.82	17.24	26.35	20.12	19.44	10.67	18.88		
HENBA1005530	166.03	123.7	313.84	376.32	270.85	265.42	127.01	98.49	128.67		
HENBA1005538	4.22	2.81	4.67	3.91	2.23	1.75	0.82	1.49	1.34	*	-
HENBA1005539	7.37	8.42	11.33	9.58	8.07	5.43	8.22	7.08	3.71		
HENBA1005545	6.57	6.11	9.83	3.2	2.97	2.68	4.51	2.07	2.38	*	*
HENBA1005548	9.87	15.77	17.38	7.24	13.44	14.5	10.18	7.2	5.13		
HENBA1005552	51.47	60.33	76.88	44.85	67.33	45.49	61.05	38.5	41.19		
HENBA1005558	1303.4	1895.5	2079.7	1883.4	2100.7	1420	1658.2	1099.2	844.47		
HENBA1005568	18.05	24.94	43.8	19.93	30.38	19.35	24.29	10.08	5.54		
HENBA1005570	9.18	10.68	10.58	6.59	4.61	5.33	3.91	1.39	3.62	**	**

22 / 157

【0846】

【表375】

(1243)

特開2002-191363

2483

2484

MEMBA1005676	3.76	5.34	5.63	3.64	6.16	2.3	3.82	3.83	3.35
MEMBA1005677	3.45	4.03	5.12	2.3	2.95	1.9	2.63	1.06	1.56
MEMBA1005681	17.34	16.11	23.24	3.88	10.8	7.1	4.84	4.82	3.74
MEMBA1005682	240.9	366.29	385.46	317.4	608.67	348.15	243.13	155.98	318.09
MEMBA1005683	56.28	75.41	54.74	38.66	63.45	30.25	44.93	12.37	20.67
MEMBA1005688	48.19	61.31	71.13	27.32	36.77	18.93	99.81	56.7	45.14
MEMBA1005693	8.48	13.58	28.24	14.33	18.97	7.86	11.3	3.69	6.87
MEMBA1005695	5.4	4.89	14.77	3.77	3.53	5.64	3.73	2.81	4.24
MEMBA1005697	4.61	6.07	5.65	4.64	29.78	7.83	6.41	5.62	16.69
MEMBA1005698	10.67	19.2	23.77	3.88	10.94	4.17	5.64	1.39	3.32
MEMBA1005699	15.17	18.03	17.77	3.18	7.05	3.09	3.33	3.84	3.16
MEMBA1005701	3.29	3.65	8.89	1.88	4.28	2.06	3.75	2.04	3.26
MEMBA1005721	2.8	5.27	13.13	1.83	2.71	2.81	4.45	1.3	2.19
MEMBA1005727	28.17	45.68	60.66	11.35	32.13	16.73	25.61	10.31	7.66
MEMBA1005728	4.16	6.62	18.11	5.71	6.97	5.83	6.46	1.4	3.09
MEMBA1005731	2.97	2.99	6.58	2.7	2.52	4.58	1.93	1.12	1.83
MEMBA1005732	8.76	7.76	13.3	5.8	4.94	4.76	5.21	2.6	5.1
MEMBA1005734	13.26	27.71	30.18	8.65	7.52	11.69	10.2	6.61	5.28
MEMBA1005762	5.49	5.63	8.69	1.07	4.9	3.69	4.47	3.62	2.83
MEMBA1005769	4.1	8.92	17.42	4.61	5.06	7.38	8.37	3.66	4.18
MEMBA1005770	19.65	25.67	58.84	10.77	25.74	28.55	25.33	19.21	13.78
MEMBA1005771	3.25	3.78	12.85	4.13	4.03	2.88	2.63	0.58	1.69
MEMBA1005779	30.48	22.92	60.53	11.99	24.03	21.81	24.62	12.99	24.28
MEMBA1005780	5.31	5.25	11.55	7.98	6.87	8.25	4.31	2.95	4.88
MEMBA1005785	10.54	10.31	19.75	6.26	14.08	11.68	12.25	7.49	2.77
MEMBA1005793	3.05	4.09	5.98	5.76	3.35	4.26	2.89	1.71	1.85
MEMBA1005799	61.55	73.56	131.24	173.55	121.01	131.74	58.39	28.63	52.24
MEMBA1005793	8.7	6.7	19.33	15.32	12.15	10.29	11.29	5.53	9.29
MEMBA1005795	10.69	16.62	28.52	12.65	15.48	9.89	13.24	7.86	8.73
MEMBA1005712	14.41	9.07	21.89	4.62	12.08	8.24	12.54	7.16	9.37
MEMBA1005717	9.58	15.11	20.28	15.34	17.37	18.39	10.48	4.92	9.53
MEMBA1005718	14.72	17.48	32.28	23.31	19	19.82	12.33	8.89	13.45
MEMBA1005721	7.61	8.19	18.43	10.15	9	9.16	5.24	3.31	7.52
MEMBA1005722	8.98	8.38	15.66	10.88	5.16	8.22	6.16	2.81	4.2
MEMBA1005724	3.21	3.63	9.19	3.08	3.41	4.96	4.32	2.79	3.41
MEMBA1005732	26.35	14.39	37.85	48.13	33.11	35.41	27.65	13.58	24.67
MEMBA1005737	10.7	10.58	19.74	12.96	11.14	14.53	14.77	5.18	11.46
MEMBA1005742	24.18	21.22	31.29	17.65	22.38	19.25	19.89	11.57	20.16
MEMBA1005745	16.18	18.76	28.18	19.8	21.24	20.22	17.64	16.51	21.84
MEMBA1005747	4.59	6.97	9.33	5.84	4.66	4.23	3.4	2.42	2.9
MEMBA1005749	4.06	4.65	5.69	4.59	4.51	2.42	4.14	4.85	3.14
MEMBA1005755	6.51	5.76	10.77	4.34	2.86	3.2	3.26	2.6	1.41
MEMBA1005760	8.14	6.09	21.23	11.82	15.12	18.05	5.98	3.99	9.09
MEMBA1005765	16.17	18.36	30.68	12.16	17.74	19.6	18.91	12.07	18.52
MEMBA1005766	4.59	6.7	11.4	6.4	6.48	7.14	3.63	2.36	6.46
MEMBA1005780	66.74	68.34	50.49	70.67	83.79	79.9	51.01	23.87	50.17
MEMBA1005795	12.49	13.63	22.28	21.84	27.08	17.3	17.89	8.53	18.83
MEMBA1005809	27.58	26.57	37.75	10.62	11.75	10.76	7.41	20.02	13.83
MEMBA1005813	11.01	13.12	21.07	10.05	5.62	7.48	3.76	4.75	2.72
MEMBA1005815	13.68	21.48	26.3	18.7	15.38	15.64	15.88	14.09	23.04
MEMBA1005822	2.32	4.48	5.72	1.33	1.78	1.99	1.83	4.15	2.65
MEMBA1005829	11.83	17.29	26.15	14.23	14.49	18.69	18.12	8.88	16.12
MEMBA1005833	7.51	11.08	18.85	8.38	8.46	9.74	7.13	4.47	9.24
MEMBA1005834	2.53	4.28	9.15	2.39	5.2	3.43	3.05	1.67	2.22
MEMBA1005844	5.49	8.18	12.47	6.98	9.74	6.87	6.18	2.19	5.4
MEMBA1005852	8.05	12.52	17.23	5.22	7.99	5.69	7.73	5.09	7.72
MEMBA1005853	12.09	27.28	20.82	6.51	8.22	8.09	4.4	8.87	5.87
MEMBA1005878	18.48	26.29	29.76	6.78	14.33	12.64	11.87	13.79	15.84
MEMBA1005883	10.6	8.45	21.85	5.18	6.88	6.16	3.04	2.65	2.62
MEMBA1005884	18.4	17.45	38.78	13.72	28.17	26.47	16.14	9.17	16.88
MEMBA1005891	14.35	23.5	29.7	12.98	17.06	15.29	15.47	12.21	17.4
MEMBA1005894	20.78	25	45.44	20.3	23.69	18.03	21.15	9.72	19.92
MEMBA1005898	6.21	7.33	9.72	7.72	7.65	5.64	3.88	1.69	3.35
MEMBA1005902	62.63	64.98	95.93	140.97	103.88	141.36	40.77	40.68	57.15

23 / 157

【0847】

【表376】

2485

2486

HENBA1005907	11.51	24.95	21.58	5.18	12.98	7.51	14.52	20.83	32.4				
HENBA1005909	12.54	10.65	15.16	9.37	9.7	8.92	5.33	5.2	5.61				
HENBA1005911	4.25	2.94	8.89	4.51	4.07	3.47	3.19	0.81	2.14				
HENBA1005912	25.05	30.41	48.44	55.56	35.89	54.18	35.55	16.89	26.27				
HENBA1005913	8.07	12.88	17.18	3.35	11.55	8.12	4.58	4.08	5.57				
HENBA1005921	8.39	17.95	29.27	13.5	16.47	18.42	20.02	11.79	13.13				
HENBA1005922	10.23	12.69	22.71	8.62	19.47	16.43	10.1	8.02	9.38				
HENBA1005929	14.03	14.99	35	18.43	15.07	16.08	12.42	10.95	15.82				
HENBA1005931	6.92	7.45	8.96	7.96	7.11	4.38	5.8	5.53	5.56				
HENBA1005934	9.15	11.61	14.52	8.47	9.42	7.17	8.08	6.07	3.78				
HENBA1005945	13.18	17.17	24.76	14.61	10.38	10.87	14.78	9.7	14.04				
HENBA1005962	8.03	8.91	21.61	8.95	9.17	9.16	8.64	3.56	9.28				
HENBA1005963	7.03	8.04	15.83	2.68	5.58	6.69	5.62	2.88	7				
HENBA1005990	1.77	5.11	10.33	2.7	4.79	2.9	2.4	1.78	3.08				
HENBA1005991	84.98	101.38	91.59	21.08	89.91	49.01	45.05	21.27	28.13				
HENBA1005993	10.77	8.18	14.65	9.74	9.96	11.62	8.75	7.35	11.3				
HENBA1006002	4.73	10.03	9.92	4.44	5.47	3.19	2.29	3.3	1.98				
HENBA1006005	11.33	12.26	21.89	15.72	14.29	10.83	11.77	7.68	11.25				
HENBA1006011	7.74	6.02	9.68	4.17	3.85	3.85	4.89	1.78	2.14				
HENBA1006013	15.75	24.43	27.02	14.33	11.65	10.41	12.44	6.78	8.78				
HENBA1006018	12.67	16.72	32.68	4.67	8.33	5.18	11.34	2.62	5.68				
HENBA1006019	39.95	43.51	94.43	41.01	47.95	33.8	48.85	26.74	35.48				
HENBA1006021	100.18	84.95	188.84	161.47	154.88	173.68	76.07	51.55	67.34				
HENBA1006022	4.41	4.65	8.97	5.38	4.93	5.29	5.93	2.38	6.8				
HENBA1006031	54.5	65.09	83.41	23.76	30.34	33.23	14.55	24.74	14.18				
HENBA1006035	26.81	28.57	28.41	13.64	17.75	18.2	23.49	18.85	24.27				
HENBA1006036	30.41	35.58	51.17	28.85	21.84	21.47	22.61	12.69	16.92				
HENBA1006042	14.78	15.89	80.2	21.62	12.62	17.35	12.05	7.08	10.83				
HENBA1006044	6.37	6.38	15.67	2.89	5.13	3.3	6.8	2.77	4.16				
HENBA1006045	5.05	5.88	16.26	5.37	6.61	3.88	4.28	3.34	4.33				
HENBA1006048	5.68	8.21	17.12	5.57	7.7	7.23	7.81	2.07	6.93				
HENBA1006053	9.91	12.14	22.17	11.18	10.11	10.25	6.6	4.29	8.53				
HENBA1006056	18.11	16.89	29.58	15.19	13.98	12.78	12.35	11.58	12.66				
HENBA1006058	12.02	10.46	14.5	9.27	7.95	5.43	9.22	7.8	6.16				
HENBA1006083	115.82	107.87	141.14	181.05	128.88	130.01	94.43	64.84	83.98				
HENBA1006087	14.48	16.19	18.97	8.28	13.91	5.01	10.48	4.28	4.88				
HENBA1006081	3.95	8.3	15.7	2.09	6.94	3.38	3.92	1.35	3.37				
HENBA1006089	4.09	6.89	17.6	2.93	4.65	1.68	3.54	1.57	3.85				
HENBA1006090	16.51	33.88	39.21	14.18	23.81	18.25	18.94	5.99	5.67				
HENBA1006091	6.4	6.84	10.5	5.04	5.34	5.02	4.02	3.3	3.85				
HENBA1006093	24.23	35.95	33.59	5.11	7.78	13.97	5.79	10.1	7.62				
HENBA1006098	23.7	22.12	46.77	18.77	21.87	13.39	21.08	17.59	21.54				
HENBA1006100	15.19	17.46	28.4	13.14	14.55	10.6	18.9	11.15	17.08				
HENBA1006108	9.23	5.57	12.26	4.66	7.02	4.99	4.59	2.55	4.37				
HENBA1006114	6.24	4.2	16.8	2.53	7.18	4.48	5.37	0.85	3.06				
HENBA1006121	4.96	7.47	17.94	3.37	8.22	2.88	5.11	1.02	2.9				
HENBA1006124	23.37	10.77	43.63	10.18	13.13	12.16	9.91	5.78	6.28				
HENBA1006125	5.3	5.09	9.35	4.55	4.9	6.58	2.71	1.37	4.21				
HENBA1006190	81.41	100.18	218.91	239.88	178.95	190.17	100.6	68.45	89.03				
HENBA1006198	16.78	21.47	42.54	20.91	18.85	14.71	18.67	11.93	13.92				
HENBA1006142	5.26	5.14	9.74	3.49	5.53	4.08	2.75	2.92	2.44				
HENBA1006150	20.94	16.77	47.47	16.05	12.94	20.13	22.73	16.22	18.83				
HENBA1006151	3.81	3.82	12.07	4.08	2.71	3.79	4.65	1.5	3.25				
HENBA1006155	51.87	36.97	81.85	14.8	33.19	24.55	29.88	21.38	8.77				
HENBA1006158	7.8	5.18	16.23	4	7.84	5.44	10.27	1.55	3.09				
HENBA1006164	7.41	4.45	10.51	11.19	6.82	6.2	5.41	2.39	2.71				
HENBA1006171	11.38	13.76	11.95	10	8.74	12.84	7.5	4.2	3.45				
HENBA1006173	5.77	5.17	5.55	5.75	11.01	7.35	5.28	2.14	3.63				
HENBA1006176	10.21	6.39	20.68	12.38	13.62	14.28	6.26	8.81	6.78				
HENBA1006182	6.52	8.53	12.82	8.56	9.65	11.37	5.71	3.22	5.23				
HENBA1006187	4.36	5.58	6.45	4.07	4.78	4.63	3.18	1.47	2.61				
HENBA1006198	6.83	8.29	18.43	9.27	9.88	14.83	8.95	4.93	5.04				
HENBA1006213	48.5	59.16	87.78	45.37	52.9	41.81	64.89	38.82	49.81				
HENBA1006217	77.53	104.19	150.85	44.45	56.82	81.63	58.2	51.63	85.92				

24 / 157

【0848】

【表377】

2487

HEMBA1006226	60.68	89.4	128.3	98.59	21.45	56.73	34.83	42.34	47.3
HEMBA1006225	7.29	6.42	9.61	4.4	5.83	6.34	4	1.9	2.78
HEMBA1006248	8.59	14.3	12.76	17.84	10.1	10.62	14.29	11.1	13.06
HEMBA1006251	15.98	24.11	24.67	22.82	19.08	21.09	18.03	3.32	8.44
HEMBA1006252	6.29	11.35	24.58	17.83	5.75	7.22	9.32	6.16	9.75
HEMBA1006253	8.65	9.14	20.83	14.78	17.19	22.49	8.79	1.77	6.4
HEMBA1006259	11.81	21.15	28.91	16.9	10.94	16.19	15.68	14.83	17.05
HEMBA1006261	25.45	37.82	48.84	33.85	29.71	14.07	18.76	16.97	24.68
HEMBA1006268	4.53	5.62	11.14	5.03	3.83	5.91	4.8	2.63	2.52
HEMBA1006271	27.87	48.98	84.46	39.21	39.6	40.82	30.74	14.23	28.33
HEMBA1006272	3.48	5.09	12.53	3.82	7.27	10.33	8.77	2.68	5.16
HEMBA1006273	4.26	5.62	9.16	7.22	8.74	8.25	4.33	1.01	3.78
HEMBA1006275	8	7.68	13.01	10.92	8.88	11.33	8.59	5.55	8.36
HEMBA1006278	3.56	6.86	4.97	4.58	7.03	4.18	3.68	1.91	2.81
HEMBA1006283	21.81	22.15	38.14	17.11	13.97	13.64	11.04	8.64	8.53
HEMBA1006284	9.77	12.43	9.21	7.3	6.48	8.66	4.11	16.95	4.13
HEMBA1006291	7.57	5.98	8.47	4.12	5.97	5.88	4.51	2.35	4.32
HEMBA1006292	28.37	23.25	41.05	48.83	29.24	37.1	21.69	20.86	21.95
HEMBA1006293	3.75	5.49	8.43	3.08	2.38	10.01	0.91	2.39	2.11
HEMBA1006299	29.83	28.58	31.82	37.12	28.94	29.71	18.96	13.53	20.58
HEMBA1006309	12.21	13.85	18.63	10.12	12.48	15.92	14.59	3.8	13.91
HEMBA1006310	19.98	14.98	14.85	11.71	17.38	10.38	10.75	8.47	4.15
HEMBA1006311	7.24	7.84	18.35	13.11	5.74	8	7.15	4.03	4.85
HEMBA1006313	3.22	8.11	10.2	4.45	3.61	3.07	2.82	2.44	2.58
HEMBA1006318	7.65	12.92	16.52	10.38	5.11	11.89	7.9	6.94	8.74
HEMBA1006328	21.63	21.11	38.33	32.37	21.64	26.1	17.68	13.27	19.38
HEMBA1006334	3.37	3.68	14.62	4.58	3.28	2.72	2.89	4.86	1.73
HEMBA1006335	17.68	32.85	48.47	23.14	32.25	24.21	26.2	15.13	17.28
HEMBA1006344	55.44	64.87	82.83	51.59	45.13	40.79	32.94	23.43	27.33
HEMBA1006347	7.7	7.81	11.32	7.88	6.93	5.21	8.91	1.81	2.49
HEMBA1006349	5.12	7.39	13.08	11.88	7.82	10.21	7.73	3.41	9.88
HEMBA1006352	4.71	5.44	5.8	3.17	4.05	8.34	3.53	2.18	6.11
HEMBA1006357	21.84	30.15	32.51	18.24	19.4	20.03	17.68	17.67	25.75
HEMBA1006358	9.29	7.19	14.9	8.1	8.4	7.37	7.12	3.93	6.97
HEMBA1006359	5.5	5.93	13.38	4.65	8.68	6.77	4.91	2.59	3.15
HEMBA1006360	4.48	4.17	9.74	2.49	4.18	4.08	2.89	1.16	2.68
HEMBA1006364	4.83	5.25	14.78	2.78	4.9	3.21	2.71	2.21	3.79
HEMBA1006377	88.85	114.32	105.84	75.76	102.38	58.33	33.99	60.04	78.87
HEMBA1006380	29.03	20.23	49.45	23.08	19.66	23.19	14.88	15.28	25.75
HEMBA1006381	21.8	26.78	34.04	28.88	18.12	16.56	28.76	15.64	19.65
HEMBA1006385	26.15	27.47	48.23	21.69	21.93	26.5	17.08	11.15	28.63
HEMBA1006390	68.78	79.54	88.03	47.23	63.78	61.2	79.04	51.89	65.97
HEMBA1006391	9.83	7.64	21.2	6.37	10.85	8.66	7.1	4.25	6.68
HEMBA1006398	7.74	10.16	17.84	6.15	9.15	10.18	5.31	3.87	8.38
HEMBA1006405	17.21	30.3	38.87	11.67	17.28	9.3	9.52	5.99	10.52
HEMBA1006410	12.29	24.19	20.82	11.01	22.54	19.08	10.32	24.08	7.5
HEMBA1006416	13.77	16.34	35.1	20.2	18.08	14.96	13.83	12.25	17.9
HEMBA1006418	14.68	14.23	12.55	8.74	10.37	7.82	5.57	12.05	3.94
HEMBA1006419	39.32	28.38	45.23	21.57	20.76	20.5	33.29	23.18	35.96
HEMBA1006421	6.94	7.3	14.62	4.48	5.03	4.44	5.76	2.74	5.79
HEMBA1006424	6.59	6.16	10.85	2.39	3.13	4.03	3.8	3.29	5.21
HEMBA1006428	16.94	21.77	35.51	13.82	15.8	11.92	15.06	10.89	16.65
HEMBA1006430	21.51	23.1	52.14	21.8	26.01	14.55	25.55	15.62	24.71
HEMBA1006438	7.39	9.88	27.72	9.7	12.88	10.02	7.77	3.89	8.06
HEMBA1006445	6.97	4.62	11.24	6.9	4.18	5.84	3.07	2.65	7.57
HEMBA1006446	10.44	28.9	25.23	12.03	4.68	7.64	2.88	8.87	4.85
HEMBA1006456	20.27	20.9	27.71	19.85	18.2	23.69	16.72	10.9	15.13
HEMBA1006461	10.85	13.87	19.68	8.91	8.18	8.85	10.75	7.35	8.75
HEMBA1006467	2.98	4.33	9.77	4.94	3.17	2.58	4.08	1.78	2.14
HEMBA1006470	34.3	38.7	56.67	41.16	36.04	29.19	28.2	24.42	29.45
HEMBA1006471	8.92	14.85	23.43	12.81	12.27	40.2	16.21	5.14	6.51
HEMBA1006474	80.88	55.42	113.87	83.82	77.51	77.92	59.58	47.32	61.33
HEMBA1006476	124.18	125.67	311.61	282.64	262.79	111.47	88.12	112.31	128.83
HEMBA1006482	750.18	800.51	717.99	555.6	740.67	596.85	755.76	752.48	372.51

25 / 157

2488

【0849】

【表378】

2489

2490

HEMBA1005483	10.38	11.79	17.87	10.54	8.08	5.41	9.38	6.26	10.8				
HEMBA1005485	14.11	21.34	22.05	9.69	9.42	8.66	9.41	4.35	6.34	*	*	-	-
HEMBA1005486	25.8	38.65	46.49	15.25	21.42	18.89	28.29	12.46	17.42	*	*	-	-
HEMBA1005489	4.69	5.66	11.89	2.18	3.64	2.6	4.74	2.19	2.49				
HEMBA1005492	21.28	21.32	31.79	8.11	21.1	11.89	18.25	6.33	8.88	*	*	-	-
HEMBA1005494	3.73	5.82	13.17	4.37	5.86	3.22	4.57	1.78	4.89				
HEMBA1005497	9.48	5.39	9.72	8.41	4.7	6.22	4.84	2.41	5.11				
HEMBA1005501	70.71	74.3	109	127.89	121.18	132.43	58.47	47.55	64.22	*	*	+	-
HEMBA1005502	28	37.84	37.28	17.65	19.66	21.79	14.53	12.47	14.09	*	**	-	-
HEMBA1005507	74.35	71.35	93.66	76.72	76.18	89.01	64.44	46.82	61.01				
HEMBA1005517	10.16	13.34	25.2	8.05	9.91	8.2	9.38	8.78	9.53				
HEMBA1005521	4.71	7.89	15.95	2.17	4.99	3.48	4.73	1.47	4.08				
HEMBA1005529	20.44	40.14	54.85	11.1	18.97	9.04	8.44	3.17	5.16	*	*	-	-
HEMBA1005530	2.11	3.93	18.06	3.3	3.94	2.97	2.36	0.8	2.85				
HEMBA1005535	4.76	6.35	8.43	4.99	4.07	4.99	1.52	1.3	3.38	*	*	-	-
HEMBA1005536	7.73	9.35	15.74	6.97	6.8	8.19	6.06	4.78	7.71				
HEMBA1005540	4.89	5.07	10.1	3.8	4.1	3.21	4.5	0.67	2.67				
HEMBA1005544	9.55	9.44	12.65	5.51	8.8	3.64	6.81	6.28	5.66	*	*	-	-
HEMBA1005548	19.83	18.34	35.37	18.49	13.07	8.02	20.42	13.65	12.24				
HEMBA1005549	3.89	4.64	10.47	3.6	3.22	3.38	4.38	1.91	2.6				
HEMBA1005559	5.63	7.78	15.83	4.1	6.24	3.33	4.33	1.27	1.6				
HEMBA1005562	4.46	7.73	13.41	4.81	4.38	7.66	8.04	1.8	5.11				
HEMBA1005566	1.87	2.79	6.04	4.85	3.75	3.09	2.83	0.98	3.24				
HEMBA1005569	9.03	13.38	17.77	8.08	5.02	11.82	6.18	3.56	6.28	*	*	-	-
HEMBA1005572	4.01	5.33	6.28	3.85	4.61	4.82	3.06	1.11	1.97	*	*	-	-
HEMBA1005579	48.81	48.67	106.01	77.85	58.85	72	32.03	36.94	29.18				
HEMBA1005583	21.83	28.38	62.62	37.84	32.54	33.88	32.6	18.08	30.33				
HEMBA1005595	9.39	11.2	24.82	12.48	12.18	12.39	11.97	4.53	8.41				
HEMBA1005597	10.74	17.9	34.13	14.71	18.52	13.99	17.64	8.01	20.29				
HEMBA1005606	9.14	17.8	25.4	12.84	13.76	11.34	17.49	5.73	12.78				
HEMBA1005812	18.18	18.45	27.38	18.16	10.37	13.64	7.68	7.75	9.58	*	*	-	-
HEMBA1005817	8.08	9.55	19.24	13.65	9.22	10.71	8.25	4.71	8.01				
HEMBA1005824	62.58	68.3	115.43	55.71	47.08	62.71	45.67	24.82	44.31				
HEMBA1005831	23.12	25.87	43.15	23.48	22.27	20.06	24.55	15.53	27.45				
HEMBA1005835	7.91	10.88	14.85	9.02	10.78	9.39	11.45	8.88	11.05				
HEMBA1005839	10.04	12.39	17.07	18.68	19.19	12.58	8.27	4.47	14.67				
HEMBA1005843	1.84	4.34	8.2	4.98	3.53	3.26	2.93	0.82	1.75				
HEMBA1005848	15.63	17.03	27.49	19.15	15.29	16.87	5.29	8.82	12.26				
HEMBA1005852	28.71	34.75	50.3	32.87	21.78	18.68	16.24	13.51	23.67				
HEMBA1005853	7.58	8.61	17.15	7.18	4.77	7.22	2.33	2.95	3.74				
HEMBA1005858	19.91	14.88	48.21	18.71	18.4	21.38	11.54	8.35	7.4				
HEMBA1005859	58.61	60.78	125.49	159.5	103.62	108.54	57.82	48.19	67.43				
HEMBA1005865	2.48	4.7	11.42	4.57	4.2	3.45	4.37	1.98	4.61				
HEMBA1005868	3.78	4.89	6.54	3.83	3.74	5.58	5.07	2.85	4.33				
HEMBA1005871	28.68	48.04	31.81	11.03	10.78	7.48	20.4	8.57	17.2	**	*	-	-
HEMBA1005874	17.95	21.64	48.41	21.77	17.05	22.65	20.34	20.43	18.22				
HEMBA1005876	8.33	14.71	17.2	11.8	8.88	5.97	6.74	4.48	8.11	*	*	-	-
HEMBA1005882	6.28	7.81	8.43	5.22	3.34	3.44	5.44	2.35	3.91				
HEMBA1005888	8.05	7.99	15.78	5.58	7.11	7.53	6.82	4.88	5.53				
HEMBA1005895	11.68	16.81	23.72	8.88	10.42	13.13	10.65	9	11.41				
HEMBA1005898	12.25	15.75	12.83	8.51	9.6	8.75	4.49	2.19	4.51	*	**	-	-
HEMBA1005702	6.88	8.14	9.5	5.54	6.34	5.88	4.33	1.73	4.83	*	*	-	-
HEMBA1005707	12.13	15.02	19.11	19.48	16.83	14.84	14.25	5.83	13.01				
HEMBA1005708	7.17	12.19	21.37	12.13	8.08	8.49	3.77	4.26	7.59				
HEMBA1005709	13.09	19.04	28.5	14.85	14.05	15.88	11.87	11.01	16.42				
HEMBA1005717	5.25	5.26	12.53	4.74	4.01	3.68	7.38	2.38	6.67				
HEMBA1005724	12.11	8.3	19.78	5.53	10.32	5.8	9.44	6.24	8.18				
HEMBA1005731	7.48	7.6	18.92	6.87	8.78	6.75	5.19	3.77	3.96				
HEMBA1005737	6.06	6.19	10.33	4.37	4.14	4.69	4.38	0.91	4.4				
HEMBA1005742	8.98	8.7	24.13	7.5	8.95	8.07	10.84	6.04	17.08				
HEMBA1005743	14.91	22.68	22.54	17.15	17.28	14.05	13.93	8.87	7.89	*	*	-	-
HEMBA1005744	17.72	19.61	39.68	28.17	20.27	23.13	18.18	15.85	23.07				
HEMBA1005749	4.84	4.51	4.85	3.84	2.79	5.06	3.99	2.53	5.17				
HEMBA1005752	74.49	114.01	108.5	80.8	62.77	63.46	58.97	93.88	63.11				

26 / 157

【0850】

【表379】

2491

2492

MEMBA1006754	10.3	5.77	11.65	6.17	5.72	5.75	6.13	5.01	8.39				
MEMBA1006758	7.27	4.9	12.62	10.94	5.09	5.73	5.5	2.15	4.84				
MEMBA1006767	5.55	6.53	9.74	2.73	5.00	3.33	3.97	3.29	3.02				
MEMBA1006770	10.91	18.24	23.01	9.34	17.17	9.49	6.46	5.37	4.52				
MEMBA1006779	21.14	29.08	48.61	19.18	32.07	21.74	27.02	14.11	23.35				
MEMBA1006780	31.68	41.91	71.99	39.02	25.6	33.44	20.75	21.87	35.28				
MEMBA1006789	11.15	8.92	10.14	6.24	5.48	5.24	3.29	3.78	6.02				
MEMBA1006795	13.29	13.12	23.2	10.87	8.5	11.23	14.12	7.4	15.89				
MEMBA1006798	8.15	7.65	12.79	2.99	3.39	3.14	3.97	0.85	2.9				
MEMBA1006805	10.17	13.14	20.22	7.25	5.84	4.88	8.13	5.47	7.83				
MEMBA1006807	358.41	425.1	450.14	425.58	524.37	388.31	478.65	356.88	265.31				
MEMBA1006813	3.32	4.75	8.38	0.01	2.85	1.47	2.38	0.6	1.87				
MEMBA1006819	11.44	14.72	25.12	8.66	18.43	11.48	10.78	2.32	9.36				
MEMBA1006821	15.78	14.89	42.37	23.12	22.95	19.27	10.58	11	21.11				
MEMBA1006824	31.97	30.89	49.18	24.26	23.02	19.71	31.05	26.34	34.02				
MEMBA1006832	259.38	308.28	291.59	339.91	237.21	321.61	326.22	280.59	176.89				
MEMBA1006834	46.44	65.27	67.81	62.09	43.11	55.81	73	47.8	42.25				
MEMBA1006835	8.46	14.27	21.58	9.55	10.78	10.45	9.6	6.1	8.18				
MEMBA1006843	18.15	18.91	28.2	12.23	21.78	20.94	21.53	7.17	5.58				
MEMBA1006849	26.18	26.08	44.03	23.34	31.15	21.57	33.38	14.9	24.89				
MEMBA1006850	16.77	33.58	37.75	10.59	18.63	8	9.03	3.5	6.56				
MEMBA1006861	14.67	13.53	34.61	20.34	16.16	19.04	12.38	6.63	17.89				
MEMBA1006865	39.2	34.77	61.65	71.58	60.52	57.08	33.27	28.65	36.65				
MEMBA1006867	18.5	15.53	22.34	9.22	8.88	9.34	9.42	6.3	3				
MEMBA1006873	17.71	23.93	26.16	14.12	11.71	14.57	15.58	7.18	17.11				
MEMBA1006877	7.4	13.49	15.12	6.73	6.8	3.67	5.7	2.95	4.47				
MEMBA1006878	10.48	14.11	25.64	11.23	13.75	11.88	9.01	2.84	7.38				
MEMBA1006879	12.94	16.27	24.73	12.26	12.07	8.95	12.1	3.4	9.76				
MEMBA1006884	16.51	27.4	33.59	6.17	20.26	51.39	18.52	5.59	4.55				
MEMBA1006885	29.88	29.68	41.54	20.77	25.78	20.37	13.71	10.63	21.42				
MEMBA1006888	94.54	114.98	88.8	8.15	45.42	40.45	16.06	32.92	14.53				
MEMBA1006889	11.07	12.38	15.21	4.3	6.85	5.29	4.87	2.48	1.84				
MEMBA1006896	72.84	75.71	85.1	26.42	32.71	26.4	32.9	16.83	9.13				
MEMBA1006900	25.47	32.8	47.12	42.47	30.76	32.5	22.98	11.97	13.31				
MEMBA1006902	4.95	12.4	20.5	3.85	6.21	3.85	6.17	0.96	2.8				
MEMBA1006912	17.87	28.61	49.99	20.94	22.1	17.02	26.3	12.33	21.55				
MEMBA1006914	76.35	138.05	152.36	81.88	82.01	60.45	65.41	43.28	74.2				
MEMBA1006918	16.23	15.18	19.35	6.9	10.02	9.35	4.27	4.91	5.74				
MEMBA1006921	5.69	9.45	11.27	4.84	7.39	7.2	5.86	4.4	4.47				
MEMBA1006925	14.39	14.83	25.68	4.9	10.25	8.6	7.17	9.16	6.2				
MEMBA1006927	8.1	6.16	12.12	3.55	6.9	3.55	4.27	4.44	3.58				
MEMBA1006929	20.15	20.6	39.98	22.07	23.94	22.4	25.87	18.49	21.87				
MEMBA1006936	9.54	10.99	23.15	1.82	9.22	9.34	13.47	4.19	5.58				
MEMBA1006938	3.12	8.25	12.19	4.64	5.27	8.86	3.85	0.26	1.77				
MEMBA1006941	38.07	33.9	81.07	14.16	37.03	19.11	37.54	19.57	9.16				
MEMBA1006942	17.04	18.44	31.05	13.54	11.67	15.34	9.95	7.49	11.08				
MEMBA1006945	67.84	90.41	117.86	45.68	44.82	62.84	68.07	38.55	55.08				
MEMBA1006949	5.04	6.37	14.93	11.38	7.05	7.71	7.34	3.55	6.74				
MEMBA1006952	4.19	5.26	9.54	3.87	5.23	3.84	4.22	1.4	1.92				
MEMBA1006960	34.21	40.14	82.89	40.95	44.04	37.38	62.24	38.27	39.73				
MEMBA1006973	11.14	12.47	24.84	14.07	12.49	9.94	10.47	6.46	8.01				
MEMBA1006974	9.69	21.72	26.59	5.58	18.54	8	10.23	6.88	3.65				
MEMBA1006978	5.49	4.82	13.28	6.87	2.91	2.68	2.79	1.22	3.42				
MEMBA1006989	5.64	5.97	15.87	14.35	9.67	15.25	5.09	3.23	6.48				
MEMBA1006993	22.88	33.04	62.59	42.82	36.88	35.47	29.4	20.95	25.28				
MEMBA1006996	6.77	6.15	10.45	9.29	6.92	7.56	4.85	2.62	3.17				
MEMBA1007001	13.8	18.68	46.24	28.01	27.9	24.22	17.83	11.6	13.83				
MEMBA1007002	38.2	51.66	66.51	167.48	96.61	115.16	37.83	43.85	28.69				
MEMBA1007013	4.23	5.6	9.85	5.49	4.83	6.04	4.32	1.34	5.39				
MEMBA1007016	13.73	18.8	48.37	22.25	24.75	25.55	26.93	13.32	18.08				
MEMBA1007017	1.78	2.36	7.34	6.45	3.5	5.8	4.75	0.45	2.16				
MEMBA1007018	12.78	19.44	28.15	6.88	7.17	10.67	6.95	5.18	4.75				
MEMBA1007044	4.27	3.89	8.13	7.28	4.08	4.08	4.49	2.95	4.25				
MEMBA1007045	3.76	6.62	12.6	6.66	8.82	7.16	7.91	2.23	5.58				

27 / 157

【0851】

【表380】

2493

HENBA1007051	8.21	9.84	21.16	10.38	7.94	9.6	8.59	4.82	5.48
HENBA1007052	0.84	3.34	5.89	3.8	6.54	5	4.04	2.23	3.75
HENBA1007053	13.43	20.61	16.83	23.76	9.7	8.82	3.37	2.88	5.47
HENBA1007057	6.84	7.67	12.49	5.42	9.68	14.27	10.77	4.02	8.83
HENBA1007062	5.25	11.04	5.4	4.94	7.78	8.88	7.11	2.74	5.57
HENBA1007063	19.82	23.76	40.56	22.16	18.37	20.33	18.59	15.25	20.25
HENBA1007068	7.17	8.3	7.54	7.65	3.18	6.35	4.34	2.31	5.66
HENBA1007069	6.08	8.94	13.83	10.41	6.55	7.88	5.58	3.75	4.11
HENBA1007073	5	7.14	13.19	9.06	8.96	8.85	10.68	5.02	4.92
HENBA1007076	9.13	8.02	19.82	9.03	17.18	17.73	15.84	7.01	8.52
HENBA1007078	47.67	62.26	92.7	36.29	60.73	51.72	32.07	18.35	34.97
HENBA1007080	88	60.21	81.69	89.81	85.78	74.19	50.6	40.71	38.08
HENBA1007084	10.8	16.68	20.31	13.17	16.74	11.58	13.37	8.47	13.17
HENBA1007085	27.29	31.32	26.53	13.81	11.35	9.85	7.08	8.85	9.17
HENBA1007087	17.8	23.13	29.38	17.3	13.84	16.38	6.49	22.47	10.72
HENBA1007089	21.17	21.1	33.63	20.7	38.69	28.03	18.24	32.57	32.51
HENBA1007095	281.88	116.22	315.3	326.58	303	223.58	76.5	112.08	127.01
HENBA1007101	27.84	40.49	45.85	23.93	27.74	34.28	41.09	23.85	48.77
HENBA1007104	4.08	6	7.77	5.68	6.26	4.31	5.49	1.3	6.52
HENBA1007108	21.73	33.28	23.78	16.4	24.34	20.37	10.72	5.09	4.25
HENBA1007112	9.74	14.28	18.52	10.29	12.13	8.53	8.39	3.2	2.8
HENBA1007113	21.85	25.07	43.45	24.93	19.55	20.96	28.57	21.39	27.84
HENBA1007121	143.44	168.43	201.05	310.8	189.7	223.39	139.17	100.56	162.13
HENBA1007129	6.41	8.28	18.59	3.42	2.84	6.23	5.66	3.45	4.93
HENBA1007147	12.43	16.07	27.38	13.28	12.69	11.59	10.77	8.49	8.81
HENBA1007149	16.43	22.2	36.85	20.24	33.61	28.87	16.97	12.22	16.92
HENBA1007151	6.66	4.91	10.19	4.89	4.78	3.84	3.37	0.61	3.29
HENBA1007172	5.54	6.79	15.25	8.45	6.57	4.48	5.2	2.26	4.93
HENBA1007174	5.55	6.74	11.4	7.88	7.31	3.31	8.36	1.4	3.1
HENBA1007176	10.88	13.78	22.08	12.85	11.37	10.5	5.77	8.47	15.97
HENBA1007178	12.93	22.04	21.56	10.13	15.89	11.16	5.44	5.45	3.49
HENBA1007185	12.08	13.89	28.24	6.12	12.57	13.67	7.16	7.65	10.67
HENBA1007188	7.21	3.2	19.91	4.57	4.73	5.82	4.03	0.89	3.73
HENBA1007194	17.48	18.75	30.8	17.47	18.26	18.1	18.81	10.34	19.89
HENBA1007200	4.21	4.82	12.05	2.35	4.05	3.94	2.19	1.93	1.96
HENBA1007203	17.41	23.17	30.14	16.76	21.31	8.75	11.68	7.36	6.25
HENBA1007206	25.5	42.75	37.89	30.71	35.28	24.97	38.96	21.71	38.63
HENBA1007224	24.39	40.17	34.49	15.91	19.51	9.53	6.52	8.53	10.32
HENBA1007226	89.72	73.23	106.85	134.34	88.67	116.09	87.3	41.82	79.85
HENBA1007240	13.64	17.35	25.89	6.08	7.14	11.16	4.24	5.15	6.41
HENBA1007241	5.35	7.7	9.62	3.24	3.12	3.91	4.59	2.68	3.21
HENBA1007242	4.41	5.32	8.31	2.05	4.24	3.88	2.68	0.75	5.03
HENBA1007243	111.71	101.13	126.82	187.45	120.72	165.9	80.87	85.01	83.79
HENBA1007251	4.65	6.58	13.42	3.87	5.2	4.27	3.75	1.77	3.34
HENBA1007256	7.71	17.43	18.8	11.41	15.37	11.45	13.72	5.68	6.38
HENBA1007267	27.14	33.68	57.79	28.23	30.38	27.93	22.24	21.82	27.1
HENBA1007273	5.82	10.89	12.53	5.12	4.08	4.32	3.9	1.99	5.62
HENBA1007279	8.03	8.83	14.11	6.88	6.78	5.74	5.78	2.58	3.49
HENBA1007281	4.41	8.17	11.59	1.95	1.84	1.99	3.42	2.16	2.76
HENBA1007283	16.13	14.5	18.59	11.27	10.48	9.5	8.88	8.35	11.65
HENBA1007288	19	10.12	23.07	7.33	8.9	6.31	9.07	7.22	9.19
HENBA1007291	3.94	8.53	14.21	2.35	5.74	4.15	6.79	3.29	5.42
HENBA1007298	60.38	57.45	88.44	47.01	83.94	54.92	41.98	17.99	24.76
HENBA1007300	10.43	10.11	14.15	17.47	5.8	7.51	10.92	2.8	3.93
HENBA1007301	10.28	11.16	10.5	7.08	5.07	5.8	3.91	4.02	3.67
HENBA1007319	7.15	12.6	10.93	3.57	6.52	4.89	3.16	3.33	2.24
HENBA1007320	6.22	6.62	11.18	5.17	5.74	4.76	5.15	2.1	2.94
HENBA1007322	281.14	504.93	295.82	479.23	433.13	233.63	384.76	245.29	107.4
HENBA1007323	3.87	5.98	13.69	3.15	3.64	1.67	4.69	0.43	1.85
HENBA1007326	33.32	37.02	60.88	53.8	48.19	37.22	52.37	23.73	38.43
HENBA1007327	14.51	15.7	50.08	17.82	23.55	19.83	20.48	9.78	17.51
HENBA1007332	15.93	13.1	23.53	14.44	19.68	12.93	14.13	7.11	7.2
HENBA1007341	8.88	9.53	11.63	7.92	6.35	6.27	7.48	4.06	5.93
HENBA1007342	5.2	6.9	6.99	1.75	3.62	2.75	4.48	1.49	2.47

28 / 157

2494

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2496

MEMBA1007847	18.88	22.85	29.58	14.57	19.83	11.3	16.82	11.88	19.8
MEMBA1007853	5.79	10.55	11.88	3.83	9.83	8.03	10.28	4.85	8.24
MEMBA1009005	12.22	17.76	30.82	11.82	12.78	11.43	13.73	6.39	17.44
MEMBA1009060	18.71	35.13	41.01	19.83	22.41	13.75	23.85	10.67	26.11
MEMBA1009118	26.09	49.39	73.92	38.01	40.34	30.35	48.13	23.51	34.37
MEMBA1009024	16.68	21.57	43.98	24.42	20.7	22.54	19.03	14.92	22.2
MEMBA1009025	4.76	6.51	12.98	2.67	6.38	4.72	5.89	4.38	3.53
MEMBA1009030	20.29	17.68	33.24	12.16	18.77	10.61	16.77	10.43	14.37
MEMBA1009036	14.36	14.21	18.16	2.88	10.75	5.5	3.722	6.69	3.23
MEMBA1009037	11.11	18.21	12.93	4.27	8.42	4.3	6.83	2.67	4.79
MEMBA1009039	8.22	15.6	28.39	9.92	10.66	6.88	12.88	7.81	13.85
MEMBA1009044	11.18	16.44	31.14	11.2	12.12	8.02	10.52	6.21	8.17
MEMBA1009048	8.28	14.79	19.17	7.61	10.2	7.2	15.14	3.23	4.66
MEMBA1009050	12.03	11.84	30.05	14.4	13.92	13.02	11.35	6.38	12.59
MEMBA1009054	10.07	11.4	26.35	13.58	11.4	7.91	11.76	6.55	12.28
MEMBA1009055	458.88	310.48	850.13	127.99	485.16	208.47	382.27	384.83	141.27
MEMBA1009059	37.26	47.72	97.02	40.48	50.48	38.93	47.77	37.17	34.67
MEMBA1009072	84.63	93.08	172.58	108.05	149.4	148.43	84.23	55.56	75.03
MEMBA1009081	14.03	23.77	42.88	15.97	19.18	19.48	14.35	5.84	10.81
MEMBA1009083	11.03	18.35	42.87	17.47	14.88	14.23	18.14	9.46	15.73
MEMBA1009089	12.64	18.95	30.4	15.04	14.4	12.6	22.7	8.13	18.39
MEMBA1009094	10.67	10.41	27.93	13.68	21.42	16.12	14.85	8.63	10.48
MEMBA1009087	6.26	10.91	14.78	6.71	8.5	4.54	16.21	6.91	3.77
MEMBA1009089	14.89	12.94	32.83	17.41	17.02	13.25	13.89	18.91	12
MEMBA1009103	79.17	84.17	154.39	128.48	133.3	142.91	117.04	86.44	93.96
MEMBA1009108	13.5	15.97	19.47	7.19	4.59	5.62	5.12	3.42	4.88
MEMBA1009113	4.8	10.67	20.57	6.45	7.88	9.01	9.02	3.11	6.78
MEMBA1009119	3.97	10.01	14.45	4.41	4.33	3.8	9.37	6.02	5.7
MEMBA1009133	88.66	47.28	101.21	23.18	118.11	70.04	134.7	41.78	81.57
MEMBA1009134	13.38	15.77	31.72	20.8	11.19	14.5	8.59	5.64	9.53
MEMBA1009135	24.17	26.35	69.3	97.42	94.73	49.05	25.95	21.33	28.27
MEMBA1009141	7.91	11.34	20.35	10.92	8.05	11.48	11.73	5.94	8.7
MEMBA1009144	8.4	6.94	18.36	7.4	9.42	9.69	10.21	5.05	11.22
MEMBA1009147	3.5	3.8	22	5.61	6.52	8.28	7.08	1.83	3.83
MEMBA1009152	1.94	5.3	5.59	10.78	6.38	4.1	5.98	1.88	1.55
MEMBA1009154	7.08	8.64	11.01	9.8	10.82	8.45	9.52	5.05	11.09
MEMBA1009155	7.22	13.83	18.07	9.86	12.99	12.51	9.82	4.83	10.29
MEMBA1009173	36.12	47.88	77.86	40.67	28.14	43.42	25.94	23.15	34.58
MEMBA1009175	4.3	6.41	7.7	6	3.69	3.1	3.98	4.45	4.04
MEMBA1009176	23.24	24.4	50.1	18.86	26.55	28.15	17.6	14.1	16.3
MEMBA1009198	1.73	5.17	9.68	5.19	3.63	4.88	3.31	1.03	3.4
MEMBA1009203	2.42	2.22	6.16	4.71	3	3.25	2.71	1.91	2.68
MEMBA1009209	4.99	7.73	18.18	5.65	5.94	7.5	3.72	1.31	4.4
MEMBA1009212	4.68	5.59	7.88	3.89	6.07	3.08	4.71	1.76	3.05
MEMBA1009215	10.67	20.63	27.19	17.82	25.21	15.88	26.39	12.95	25.38
MEMBA1009217	25.38	37.33	52.98	16.84	15.01	12.09	8.3	23.64	16.2
MEMBA1009218	31.37	54.57	73.09	37.79	38.62	34.06	37.08	24.66	38.49
MEMBA1009226	12.28	16.29	20.13	10.11	7.88	11.16	5.4	9.82	9.11
MEMBA1009230	5.26	3.8	7.07	0.71	2.62	2.38	2.7	2.25	2.54
MEMBA1009240	5.06	6.08	8.84	4.39	4.6	5.48	8.75	2.24	3.3
MEMBA1009244	7.48	12.78	15.39	11.19	10.88	11.37	6.63	4.57	10.35
MEMBA1009260	1.27	3.14	7.4	1.44	3.59	2.75	1.47	0.82	2.71
MEMBA1009258	10.17	18.3	24.98	15.18	17.82	13.97	14.16	8.49	16.98
MEMBA1009284	26.46	35.45	52.88	26.93	18.96	21.08	16.44	18.13	29.12
MEMBA1009286	12	14.67	15.28	6.33	5.8	4.87	9.31	5.79	7.42
MEMBA1009279	8.27	9.13	14.25	3.35	4.95	7.95	4.41	2.77	4.49
MEMBA1009274	6.98	5.53	13.54	4.71	5.62	6.84	3.24	3.87	5.65
MEMBA1009276	6.09	3.14	11.83	3.71	3.43	4.43	2.41	0.94	1.41
MEMBA1009284	8.8	5.53	21.28	3.19	4.36	4.48	4.47	2.03	4.87
MEMBA1009307	8.48	14.9	24.21	13.5	12.6	8.38	3.4	3.88	7.88
MEMBA1009309	6.61	8.47	14.17	8.25	8.84	9.39	9.88	4.08	6.44
MEMBA1009312	6.97	8.62	14.28	14.78	8.04	11.17	5.38	4.07	7.05
MEMBA1009317	7.62	4.84	7.99	3.4	3.92	3.39	4.7	5.93	4.31
MEMBA1009318	6.93	8.7	14.09	8.17	6.23	9.67	7.59	3.11	8.29

29 / 157

2497

2498

HENBB1000332	4.52	4.06	10.22	2.71	3.85	3.38	4.09	2.08	4.92				
HENBB1000335	4.77	4.71	13.27	5.2	6.34	6.3	5.05	1.38	4.17				
HENBB1000338	4.81	8.32	11.37	1.79	3.32	7.88	2.23	1.53	1.7	*	*	-	-
HENBB1000337	29.33	35.43	67.58	48.11	46.55	55.37	21.92	22.71	26.93				
HENBB1000338	34.7	54.08	50.89	33.23	38.29	35.27	61.12	24.63	33.57				
HENBB1000339	10.38	11.59	21.89	13.7	11	9.9	8.4	8.43	14.47				
HENBB1000341	7.15	6.94	11.5	5.95	5.91	8.13	8.18	5.44	9.71				
HENBB1000343	22.64	22.93	36.07	24.98	20.99	25.34	24.47	20.51	25.26				
HENBB1000354	24.93	29.83	37.49	18.95	17.28	18.25	24.74	17.57	18.26	*		-	-
HENBB1000358	4.2	3.68	9.08	2.24	3.75	2.51	2.41	0	3.07				
HENBB1000369	7.77	9.59	8.28	0	3.18	0.74	4.07	9.09	8.59	**		-	-
HENBB1000373	4.9	5.8	9.83	3.73	6.87	3.02	3.88	2	4.12				
HENBB1000374	33.91	19.77	79.58	37.61	43.98	33.27	46.4	29.37	24.06				
HENBB1000376	23.81	37.48	51.78	28.25	28.92	24.66	24.75	22.13	31.78				
HENBB1000383	18.87	23.34	28.75	7.22	8.14	5.48	8.08	3.68	4.53	**	**	-	-
HENBB1000391	18.94	23.82	39.78	18.85	17.85	18.64	28.87	13.66	21.83				
HENBB1000399	7.91	9.01	13.97	2.99	3.3	3.33	3.37	1.8	1.68	*	*	-	-
HENBB1000402	5.88	7.15	13.91	3.49	5.23	2.79	8.8	2.16	2.32				
HENBB1000404	4.49	5.13	14.15	2.37	3.5	4.1	3.34	1.25	1.08				
HENBB1000407	10.22	18.01	27.48	10.21	12.59	7.8	5.48	2.69	5.88				
HENBB1000420	8.98	13.82	23.72	10.31	11.56	10.07	12.88	5.38	9.15				
HENBB1000430	87.5	75.58	120.02	34.45	32.4	42.89	83.02	74.93	33	*		-	-
HENBB1000434	34.1	37.7	48.77	30.73	22.51	21.34	32.9	22.77	33.21	*		-	-
HENBB1000438	4.47	5.36	6.73	3.08	5.98	3.03	2.33	1.63	2.04	*	**	-	-
HENBB1000441	14.04	26.42	32.37	18.04	18.64	13.8	26.09	16.54	18.05				
HENBB1000447	26.11	49.46	42.13	39.58	42.28	10.3	28.83	15.38	20.58				
HENBB1000449	4.81	5.17	18.11	3.66	5.27	3.1	5.38	1.18	8.41				
HENBB1000453	38.48	19.41	48.95	17.29	27.48	11.55	15.42	4.83	8.75	*	**	-	-
HENBB1000455	7.98	15.4	31.12	12.14	13.6	12.23	19.02	4.77	15.17				
HENBB1000472	8.9	8.85	20.38	10.03	8.41	7.69	7.24	3.85	5.6				
HENBB1000480	18.47	20.31	37.47	18.05	17.04	11.84	20.18	15.95	20.84				
HENBB1000488	38.69	18.48	47.19	21.3	19.24	16.88	39.82	23.52	22.86	**		-	-
HENBB1000487	8.37	6.26	12.27	7.11	4.47	4.08	6.03	4.45	5.18				
HENBB1000490	32.85	71.05	76.4	38.48	33.85	29.5	47.17	24.97	28.9				
HENBB1000491	10.68	13.88	21.94	11.39	11.14	7.83	9.44	3.7	10.75				
HENBB1000492	11.93	15.38	18.28	8.6	7.89	5.65	7.78	2.28	4.01	*	*	-	-
HENBB1000493	9.37	12.88	19.63	10.93	27.47	71.08	13.51	1.84	8.84				
HENBB1000510	11.39	18.53	31.92	20.35	17	14.3	15.03	12.6	21.32				
HENBB1000518	5	8.48	9.48	3.37	2.52	5.44	4.17	3.02	4.74				
HENBB1000523	5.79	7.88	16.89	3.63	4.25	4.51	8.74	3.95	5.88				
HENBB1000523	16.51	22.61	33.14	14.54	14.09	13.74	17.51	11.52	13.77				
HENBB1000530	6.45	6.62	14.38	4.67	5.23	3.03	8.95	3.14	3				
HENBB1000542	11.01	10.05	25.81	5.58	5.42	6.88	5.84	1.28	2.47				
HENBB1000550	7.6	9.67	18.88	5.6	8.7	4.85	3.88	1.58	2.78				
HENBB1000554	23.68	38.64	60.71	31.87	34.21	23.57	36.99	19.84	22.96				
HENBB1000556	5.71	9.66	13.39	4.65	5.23	10.47	3.94	4.88	5.74				
HENBB1000564	7.31	9.82	18.45	11.81	10.28	10.82	15.74	5.78	7.4				
HENBB1000567	20.81	20.17	47.53	8.75	7.89	6.38	13.88	8.67	4.74				
HENBB1000569	5.63	6.6	12.71	2.7	3.07	4.45	4.1	3.02	2.02				
HENBB1000573	32.51	39.2	72.29	48.4	30.21	37.17	37	20.09	25.68				
HENBB1000575	14.84	28.39	42.18	17.14	17.78	15.29	22.14	11.44	18.53				
HENBB1000579	12.25	14.06	28.81	6.45	8.01	11.92	12.64	5.71	5.94				
HENBB1000585	4.75	8.58	10.45	5.85	6.89	6.63	6.35	0.91	4.03				
HENBB1000588	9.22	12.19	32.5	16.92	12.87	17.45	7.51	6.24	9.01				
HENBB1000589	11.66	15.64	23.81	18.44	15.28	16.1	11.87	7.91	7.82				
HENBB1000591	12.71	17.74	34.25	24.98	19.88	19.8	12.8	9.24	10.68				
HENBB1000592	4.57	3.51	6.52	1.88	2.6	2.47	2.07	3.49	1.32	*		-	-
HENBB1000593	91.28	145.08	255.41	284.78	289.98	282.29	122.51	82.28	150.94				
HENBB1000595	28.02	34.64	53.68	11.28	18.32	41.15	22.88	29.26	17.38				
HENBB1000598	22.01	31.15	65.04	24.88	33.54	29.35	69.32	20.78	28.46				
HENBB1000611	2.41	5.2	8.28	7.8	5.78	3.85	5.15	1.88	3.88				
HENBB1000617	53.85	47.37	108.59	110.75	65.04	84.65	42.52	37.92	47.64				
HENBB1000623	31.42	31.62	78.25	50.87	28.93	49.28	29.97	27.08	61.77				
HENBB1000630	5.75	17.04	13.07	8.82	8.82	8.78	7.98	4.89	11.58				

30 / 157

2499

2500

MEMB1000831	17.85	20.22	28.17	18.29	17.71	18	15.1	13.03	22.53				
MEMB1000832	9.09	8	17.51	9.07	8.79	9.7	5.89	4.13	8.31				
MEMB1000836	2.41	3.55	6.37	2.23	5.22	2.73	5.28	1.01	4.17				
MEMB1000837	10.09	17.58	15.59	10.48	18.82	18.4	10.07	3.67	7.99				
MEMB1000838	2.99	4.17	4.52	3.32	2.17	3.48	2.49	1.75	2.69	*			
MEMB1000842	39.5	47.38	81.48	47.61	39.24	45.01	39.82	28.49	64.57				
MEMB1000843	19.65	15.32	32.08	29.95	12.4	14.81	11.1	10.59	13.67				
MEMB1000849	12.99	15.1	28.14	18	12.05	14.32	13.77	8.35	17				
MEMB1000852	14.35	8.01	17.67	7.91	13.41	10.62	9.88	8.25	11.18				
MEMB1000855	4.61	7.58	7.57	5.6	6.2	3.31	4.16	0.72	5.12				
MEMB1000856	4.18	6.24	9.15	3.47	7.38	8.78	5.21	1.62	5.11				
MEMB1000860	16.93	27.34	0.25	1.82	1.14	0.38	23.87	15.78	19.86				
MEMB1000871	38.43	48.48	92.78	44.14	31.73	40.3	25.53	28.64	41.28				
MEMB1000873	32.99	36.97	43.71	17.4	18	19.87	6.02	26.42	8.4	**	*		
MEMB1000878	47.01	69.39	88.27	46.41	43.74	44.85	52.81	42.78	53.45				
MEMB1000884	8.8	10.16	24.24	9.32	8.77	12.64	6.98	8.48	13.11				
MEMB1000892	16.03	14.55	25.59	15.28	15.92	17.55	18.09	8.5	14.57				
MEMB1000893	1.76	2.64	8.52	1.53	4.27	2.95	4.08	0.19	8.13				
MEMB1000705	14.59	17.08	31.29	17.89	15.24	18.52	13.25	7.08	17.6				
MEMB1000705	11.29	12.22	20.56	12.35	17.3	12.09	6.4	4.7	7.15	*			
MEMB1000709	45.75	82.83	118.05	70.74	40.37	58.09	48.11	45.53	67.59				
MEMB1000714	31.43	34.95	47.38	12.83	21.37	16.81	5.14	8.31	8.35	*	**		
MEMB1000725	5.12	8.87	10.38	4.23	8.1	4.35	4.23	1.68	4.98				
MEMB1000726	8.88	1.95	5.81	0.08	0.38	0.65	0.77	3.05	1.2				
MEMB1000729	6.42	9.45	14.46	7.38	5.18	5.16	5.32	4.6	6.58				
MEMB1000738	18.18	14.39	28.4	16.66	7.78	8.45	31.84	5.87	7.73				
MEMB1000749	19.05	25.67	36.38	23.07	19.8	20.09	17.18	15.08	19.35				
MEMB1000763	2.4	3.95	6.27	1.43	7.01	5.93	4.28	0.57	3.97				
MEMB1000770	32.56	40.83	87.69	48.69	28.8	49.69	33.53	25.98	52.34	*	*		
MEMB1000774	35.04	53.35	80.31	43.91	38.98	45.26	39.09	37.25	58.98				
MEMB1000777	45.62	74.38	70.43	17.25	25.78	25.38	18.75	21.39	27.43	*	*		
MEMB1000781	8.33	5.28	11.47	6.25	5.6	4	3.41	2.88	6.03				
MEMB1000788	18.37	23.49	32.23	14.46	18.9	18.37	22.44	12.27	19.69				
MEMB1000789	23.79	24.39	34.32	14.31	19.85	15.3	18.54	10.27	14.08	*	*		
MEMB1000790	5.59	8.56	17.48	2.6	5	2.29	4.28	1.79	6.06				
MEMB1000794	18.79	28.68	40.8	14.17	25.07	14.74	20.31	17.38	24.13				
MEMB1000807	12.94	8.25	2.93	1.02	2.13	2.36	2.89	4.28	6.87				
MEMB1000809	78.77	111.21	120.34	77.13	82.88	41.88	80.78	69.07	111.88				
MEMB1000810	18.52	9.56	13.14	4.85	2.89	5.55	9.37	2.32	10.71	**			
MEMB1000821	11.49	9.52	21.32	7.91	10.41	8.87	11.4	5.99	7.83				
MEMB1000822	14.47	17.81	31.19	12.64	15.48	12.05	15.29	6.81	15.5				
MEMB1000826	7.31	8.22	12.01	4.01	6.22	5.18	2.28	1.16	3.63	*			
MEMB1000827	3.32	10.97	13.01	2.13	2.92	3.29	2.14	1.1	3.35				
MEMB1000831	2844.2	3968.9	2893.4	2288.3	2955.9	3122.6	3699.5	1769.3	603.48				
MEMB1000835	65.9	87.92	134.3	89.52	82.05	71.78	48.58	48.76	78.94				
MEMB1000840	22.85	22.57	43.28	24.08	23.34	18.79	20.99	18.48	32.32				
MEMB1000848	11.78	12.27	18.41	9.27	11.15	6.47	11.75	8.28	10.74				
MEMB1000852	6.96	7.13	13.81	5.13	5.29	6.06	3.72	3.05	2				
MEMB1000857	14.9	18.2	33.57	15.67	15.76	14.92	14.34	8.21	17.14				
MEMB1000858	18.62	21.62	40.52	17.01	24.14	29.4	19.12	14.81	25.58				
MEMB1000867	3.35	5.16	11.87	1.01	3.43	0.49	5.8	0.59	7.29				
MEMB1000870	5.48	10.65	17.84	3.33	10.48	4.74	3.16	0.41	3.45				
MEMB1000875	9.87	10.87	29.49	19.7	15.97	14.54	9.25	6.54	15.24				
MEMB1000881	16.87	17.18	33.12	17.09	13.5	10.07	12.12	11.55	18.55				
MEMB1000883	6.22	6.79	8.86	2.81	2.5	1.97	4.54	1.83	3.25	**	*		
MEMB1000887	5.98	5.57	6.2	3.69	2.17	1.88	2.97	1.27	2.62	**	**		
MEMB1000888	3.12	5.25	10.45	4.39	3.58	2.35	3.24	2.38	1.97				
MEMB1000890	18.08	35.88	23.82	8.78	18.94	41.91	13.4	7.59	9.49				
MEMB1000893	34.65	25.07	51.31	20.43	19.67	19.95	21.33	13.24	12.88				
MEMB1000900	14.11	15.32	20.53	11.04	14.54	9.38	11.58	4.15	3.3	*			
MEMB1000905	8.7	11.8	4.84	3.81	3.66	3.74	10.34	4.58	8.59				
MEMB1000908	18.34	11.07	11.18	10.97	10.82	24.1	19.82	7.15	8.29				
MEMB1000910	14.49	13.52	29.52	12.54	12.67	12.24	17.07	12.01	16.77				
MEMB1000913	62.28	41.72	87.34	40.18	51.04	32.05	56.11	40.33	43.91				

31 / 157

2501

2502

HENB81000916	28.08	60.99	57.47	25.29	36.05	22.49	27.81	17.27	27.94
HENB81000917	5.59	5.32	12.64	3.03	4.44	3.24	2.98	0.58	2.57
HENB81000927	17.8	20.89	49.41	20.99	18.29	12.76	18.78	15.67	18.4
HENB81000932	14.04	31.9	58.79	23.64	30.24	22.56	24.51	13.79	19.35
HENB81000933	73.2	82.97	106.78	17.69	17.17	51.59	30.19	46.68	77.12
HENB81000936	28.84	37.39	39.34	11.49	17.42	18.08	9.48	19.76	21.12
HENB81000939	19.01	16.47	24.62	8.78	10.18	11.8	13.33	5.75	13.18
HENB81000941	55.55	67.85	98.78	34.29	51.94	34.02	64.38	50.22	47.74
HENB81000947	13.1	15.93	33.38	11.69	15.05	7.82	14.81	8.68	11.7
HENB81000954	9.7	14.84	29.47	10.7	10.34	11.24	13.58	8.18	14.09
HENB81000959	5.31	7.05	18.81	1.63	2.59	1.53	2.83	0	2.89
HENB81000973	7.09	9.47	31.83	11.47	11.07	10.63	11.83	4.69	9.55
HENB81000975	3.47	3.38	0.39	0.25	0.11	0.78	2.84	0.48	3.5
HENB81000981	13.64	11.74	28.6	8.77	7.91	9.41	10.88	6.85	13.87
HENB81000985	21.62	19.08	51.64	22.07	25.82	18.12	29.3	14.45	17.75
HENB81000991	46.57	60.97	104.38	49.12	52.31	37.89	58.02	46.82	42.32
HENB81000996	7.55	10.13	19.44	15.43	7.58	7.76	25.49	7.93	12.48
HENB81001000	8.88	10.27	20.46	13.94	12.61	10.4	13.32	7.07	9.02
HENB81001004	4.52	20.53	22.05	7.15	7.85	4.71	7.51	4.91	9.37
HENB81001008	27.45	57.49	106.28	36.82	28.09	41.72	54.86	25.88	43.61
HENB81001011	13.02	18.59	22.13	8.85	6.12	13.6	9.78	8.17	13.65
HENB81001014	4.03	5.88	7.48	5.94	4.03	6.17	3.85	1.78	2.98
HENB81001020	38.48	45.64	94.13	62.12	42.15	44.7	42.48	28.95	25.72
HENB81001024	14.35	17.88	33.12	21.84	15.88	15.33	12.31	9.64	11.76
HENB81001026	36.33	37.82	64.87	35.84	31.03	31.81	34.99	27.91	28.23
HENB81001037	29.37	38.91	99.73	42.44	38.61	41.98	49.51	23.14	48.42
HENB81001042	49.43	19.68	79.17	12.87	30.43	38.57	43.54	27.06	24.73
HENB81001046	1.02	3.17	4.37	2.13	2.63	1.33	3.19	0	2.81
HENB81001047	3.32	3.99	7.05	5.87	5.14	9.57	4.92	2.39	5.02
HENB81001048	14.93	18.87	36.47	38.03	15.11	21.97	9.91	8.61	13.37
HENB81001051	16.67	10.51	26.21	24.66	18.42	23.37	18.85	8.02	15.72
HENB81001056	5.76	6.07	15.19	9.62	10.04	7.15	7.81	2.99	9.72
HENB81001058	13.02	13.37	20.82	14.04	12.97	13.6	13.64	8.33	17.02
HENB81001060	8.48	6.79	11.56	3.09	7.6	8.52	5.42	2.73	5.08
HENB81001063	7.02	9.34	7.27	9.14	6.79	5.01	8.52	5.39	11.18
HENB81001068	12.68	13.51	18.25	11.13	9.67	11.92	8.97	9.12	12.67
HENB81001082	19.71	21.28	25.07	28.2	26.41	15.74	14.59	12.53	27.85
HENB81001095	9.14	8.65	18.5	8.62	8.69	8.93	5.69	6.23	9.72
HENB81001096	44.08	45.29	88.45	46.23	37.21	42.49	31	20.55	25.13
HENB81001101	14.24	13.37	19.97	9.53	14.04	14.7	13.73	9.28	15.99
HENB81001102	5.29	8.01	11.42	4.54	5.69	5.17	5.82	3.61	6.55
HENB81001104	38.08	39.86	49.87	41.44	38.7	31.13	23.82	16.12	29.93
HENB81001105	44.24	54.5	42.57	26.78	34.74	25.98	40.78	32.97	47.59
HENB81001112	7.32	12.27	12.45	6.87	5.72	6.06	2.52	6.54	5.01
HENB81001113	12.36	16.73	27.67	10.87	11.17	11.23	11.03	6.35	13.81
HENB81001114	10.5	25.04	16.53	8.31	9.62	7.81	7.26	6.53	13.76
HENB81001115	5.38	5.54	12.69	5.57	5.33	6.84	5.79	8.01	10.83
HENB81001117	24	32.48	48.34	30.8	28.94	25.33	34	33.8	48.28
HENB81001119	6.59	8.79	10.75	5.08	5.94	5.54	5.24	3.59	5.78
HENB81001126	22.85	31.95	38.8	19.87	19.28	19.37	24.09	9.52	28.55
HENB81001133	2.49	4.42	8.71	2.43	5.35	3.57	4.22	0.99	2.59
HENB81001137	7.73	11.05	0	0.73	0.56	0.68	7.84	6.25	16.01
HENB81001142	3.94	3.85	8.48	2.73	4.14	2.88	2.5	3.32	8.04
HENB81001145	7.24	10.68	17.23	5.39	7.38	5.75	4.34	3.98	7.6
HENB81001151	3.26	5.63	5.67	3.5	3.71	2.34	0.85	2.94	4.06
HENB81001153	3.1	3.77	9.68	1.47	2.32	1.7	2.21	0	1.53
HENB81001158	5.14	4.68	9.84	3.27	3.84	3.55	2.97	1.38	2.98
HENB81001169	31.33	46.44	55.42	27.9	34.88	25.03	41.78	26.83	37.34
HENB81001170	8.62	15.69	19.05	7.59	11.25	8.63	12.14	5.58	9.38
HENB81001175	2.65	3.14	5.1	4.52	2.61	4.91	2.72	1.85	6.12
HENB81001177	5.44	6.28	5.09	1.84	3.43	6.94	2.77	2.32	6.91
HENB81001182	18.97	18.18	22.47	9.28	11.35	10.35	7.83	8.39	8.5
HENB81001192	9.02	9.64	12.65	5.82	4.73	5.18	6.79	2.78	10.81
HENB81001199	4.71	4.4	13.79	2.25	2.76	5.59	3.91	3.02	2.96

32 / 157

【0856】

【表385】

(1253)

特開2002-191363

2503

2504

HENBB1001200	13.2	15.34	26.9	9.43	12.58	11.64	13.65	8.01	17.32				
HENBB1001201	18.63	26.31	21.34	16.48	23.59	8.33	19.57	7.02	6.33				
HENBB1001202	6.61	9.43	16.76	4.47	13.17	7.28	9.02	4.73	7.13				
HENBB1001203	2.22	6.57	9.4	5.43	2.97	2.71	2.73	1.63	4.83				
HENBB1001204	9.7	12.03	15.37	9.6	10.02	9.52	8.36	4.91	10.29				
HENBB1001205	10.61	10.25	17.62	6.77	6.49	7.62	10.25	4.74	16.66				
HENBB1001206	6	5.57	10.2	5.12	3.8	3.11	4.88	3.23	6.06				
HENBB1001207	26.87	26.88	47.38	27.34	25.88	18.24	28	17.58	20.48				
HENBB1001208	18.71	16.42	17.84	9.84	8.91	9.28	7.94	2.83	4.83	**	**	-	-
HENBB1001209	17.23	21.95	42.64	15.78	29.28	10.72	28.16	18.31	25.4				
HENBB1001210	6.94	12.5	13.24	4.58	8.31	8.14	5.53	1.54	8.42				
HENBB1001211	4.87	3.91	1.57	0	1.12	0.17	1.32	0.88	4.29	*		-	
HENBB1001212	5.76	2.45	7.88	3.45	3.48	1.87	2.9	1.6	4.06				
HENBB1001213	16.88	13.35	16.65	9.1	8.18	9.85	11.54	6.31	10.57	**	*	-	-
HENBB1001214	35.65	49.79	60.5	22.41	25.43	27.28	45.78	28.88	37.22				
HENBB1001215	22.08	31.03	58.18	21.17	28.84	20.21	27.38	14.91	24.06				
HENBB1001216	28.9	34.49	70.42	28.32	28.95	21.57	41.19	24.41	34.07				
HENBB1001217	7.43	6.21	17.88	4.5	8.82	3.53	4.22	0.83	3.85				
HENBB1001218	69.18	45.43	92.09	93.11	71.97	69.72	54.64	34.46	57				
HENBB1001219	184.19	98.07	152.38	13.92	33.06	57.1	34.39	45.85	70.66	*	*	-	-
HENBB1001220	34.55	33.29	61.38	34.34	20.46	23.71	30.23	24.87	29.48				
HENBB1001221	9.97	10.71	16.81	7.88	6.97	9.33	8.95	7.28	10.69				
HENBB1001222	3.47	3.55	8.83	3.1	1.84	1.48	2.02	1.47	1.7				
HENBB1001223	16.95	79.12	35.23	25.95	31.84	28.36	30.82	15.48	12.44				
HENBB1001224	22.37	27.51	58.3	21.33	26.2	17.4	27.38	15.96	28				
HENBB1001225	4.47	6.46	16.08	5.02	4.85	3	3.41	2.52	3.43				
HENBB1001226	300.12	442.69	478.77	284.44	431.41	281.22	536.94	211.14	291.4				
HENBB1001227	11.76	27.27	23.84	14.94	18.67	41.4	14.69	12.85	20.6				
HENBB1001228	14.9	20.75	33.78	16.74	10.48	13.15	15.46	13.18	16.41				
HENBB1001229	20.54	137.19	27.35	22.15	23.27	12.7	29.77	14.5	13.73				
HENBB1001230	6.45	11.88	16.83	9.21	7.91	6.84	9.36	6.15	7.34				
HENBB1001231	4.17	4.2	11.89	3.59	3.25	2.84	5.12	2.68	3.24				
HENBB1001232	58.08	165.69	115.47	28.35	68.22	33.54	41.88	11.39	13.42	*		-	
HENBB1001233	7.12	11.9	25.13	9.45	10.78	5.58	7.57	2.53	6.72				
HENBB1001234	60.43	105.03	131	68.61	74.76	59.35	124.09	50.05	41.99				
HENBB1001235	0.63	0.14	1.28	1.92	0.74	0.41	1.38	0	1.24				
HENBB1001236	19.31	21.8	38.53	21.67	17.28	16.88	20.3	14.12	28.23				
HENBB1001237	6.1	5.5	14.88	8.52	5.57	3.43	4.24	2.59	4.91				
HENBB1001238	11.08	7.69	21.58	3.98	10.13	4.45	6.01	7.37	4.42				
HENBB1001239	14.51	8.09	19.68	7.72	9.53	5.04	11.31	3.15	5.25				
HENBB1001240	2.18	3.12	12.35	0.43	2.84	1.52	3.82	0.54	2.87				
HENBB1001241	62.05	108.41	129.68	78.51	93.58	81.19	64.58	43.45	65.69				
HENBB1001242	7.52	9.31	26.43	19.44	18.12	16.4	11.29	5.29	11.49				
HENBB1001243	8.27	9.27	21.55	13.04	8.88	12.02	9.82	4.71	9.74				
HENBB1001244	11.82	15.33	29.77	17.82	9.11	9.15	17.69	10.49	12.45				
HENBB1001245	34.9	33.46	79.88	39.05	29.8	26.79	45.62	23.04	29.37				
HENBB1001246	13.59	34.47	15.18	9.61	17.82	9.58	10.35	11.38	4.3				
HENBB1001247	6.78	6.69	14.67	4.08	3.81	3.5	3.91	2.87	4.28				
HENBB1001248	7.8	18.4	21.38	7.98	16.58	12.19	13.82	2.72	8.14				
HENBB1001249	64.98	90.97	10.61	0.27	3.72	0.15	67.41	60.83	3.42				
HENBB1001250	12.67	18.5	30.58	17.22	19.87	15.48	19.81	9.85	12.99				
HENBB1001251	1.29	2.69	4.85	6.43	2.24	3.63	0.47	1.11	0.58				
HENBB1001252	7.9	11.42	30.12	18.8	14.97	11.43	10.87	6.27	10.35				
HENBB1001253	2.04	3.94	3.81	2.8	2.74	2.85	1.42	0.21	2.14				
HENBB1001254	4.48	4.13	8.85	5.53	3.83	5.8	1.34	2.67	1.9				
HENBB1001255	11.47	23.84	24.88	19.26	14.85	15.61	11.95	7.31	16.96				
HENBB1001256	5.76	4.41	16.42	7.89	11.28	6.94	5.16	2.14	5.02				
HENBB1001257	2.91	4.4	4.11	0.76	0	0.58	5.88	1.84	0.07	**		-	
HENBB1001258	14.06	23.97	51.22	27.45	25.66	29	18.89	8.5	8.91				
HENBB1001259	6.88	11.32	20.01	14.76	14.71	20	10.35	8.2	12.32				
HENBB1001260	18.71	17.84	33.05	24.24	11.69	16.55	21.47	12.89	21.69				
HENBB1001261	19.58	21.95	43.4	20.44	16.81	19.91	22.72	13.58	24.01				
HENBB1001262	35.58	35.95	71.7	35.81	35.7	39.37	42.62	30.92	51.16				
HENBB1001263	4.44	10.09	10.48	10.32	5.73	8.72	4.72	2.95	5.93				

33 / 157

【0857】

【表386】

2505

MEMB1001456	8.69	12.18	21.38	34.84	17.29	10.08	13.2	9.97	12.41
MEMB1001461	19.67	23.1	39.66	20.5	28.42	24.75	17.54	14.4	25.15
MEMB1001463	19.58	21.33	35.61	31.93	33.77	43.28	30.72	20.88	36.42
MEMB1001464	28.01	27.99	31	14.82	14.82	16.04	16.45	13.58	23.87
MEMB1001466	16.16	16.92	27.63	19.85	13.38	15.95	13.7	10.31	15.02
MEMB1001482	17.23	20.81	35.32	19.01	19	19.25	19.83	14.79	21.88
MEMB1001500	41.82	45.64	68.26	37.83	38.48	37.77	46.2	37.01	39.45
MEMB1001505	20.94	22.59	32.35	28.78	22.08	24.15	26.64	17.19	32.69
MEMB1001521	8.53	7.82	10.6	8.16	6.3	6.64	6.41	3.73	7.14
MEMB1001527	2.99	3.88	4.38	4.6	7.15	3.37	3	1.09	4.09
MEMB1001530	3.48	3.01	4.27	1.8	4.75	3.13	7.05	4.01	3.15
MEMB1001531	36.17	50.68	53.68	24.43	15.05	19.88	14.37	19.2	25.31
MEMB1001532	17.53	24.2	32.45	18.39	15.75	13.5	15.22	13.76	17.31
MEMB1001536	12.38	16.91	23.89	12.47	13.36	10.82	9.04	12.27	13.78
MEMB1001538	66.38	84.8	143.72	91.2	78.03	69.03	101.64	71.92	118.47
MEMB1001537	11.79	11.39	16.93	9.29	11.02	10.91	6.73	5.87	10.85
MEMB1001542	14.98	22.23	29.71	13.95	16.64	16.35	12.71	5.92	15.48
MEMB1001543	23.68	28.65	30.69	20.28	24.21	31.58	17.34	19.68	58.07
MEMB1001547	23.94	50.04	39.45	14.74	15.27	12.72	12.08	2.84	4.52
MEMB1001548	3.11	4.4	5.6	1.9	1.94	2.29	3.65	4.19	3.74
MEMB1001551	3.8	7.07	10.09	4.84	4.24	4.26	3.14	1.85	6.18
MEMB1001555	8.19	9.59	15.51	4.72	6.09	5.64	3.45	5.22	5.59
MEMB1001562	17.05	18.48	38.18	19.29	13.38	17.57	18.22	12.15	22.52
MEMB1001564	38.37	50	64.37	49.41	38.27	48.75	40.56	25.48	48.57
MEMB1001565	18.11	21.08	35.2	23.9	18.29	10.69	18.09	9.8	15.5
MEMB1001566	18.83	22.88	28.98	20.6	18.61	17.83	18.18	11.82	14.18
MEMB1001572	3.68	5.99	8.15	7.26	5.87	2.52	10.81	4.11	5.2
MEMB1001585	31.65	42.11	57.35	32.5	28.22	32.05	31.45	26.22	33.57
MEMB1001585	19.82	27.08	60.2	25.12	17.34	21.21	21.42	13.56	26.31
MEMB1001588	3.6	1.4	8.8	2.53	0.56	1.61	2.28	0.38	3.78
MEMB1001595	18.38	11.37	25.05	8.89	10.88	9.92	9.72	7.65	13.65
MEMB1001595	50.23	70.48	94.67	85.43	78.99	83.08	60.67	47.9	67.78
MEMB1001599	12.08	19.55	22.95	3.76	11.11	7.77	4.14	3.06	5.8
MEMB1001603	15.68	23.5	36.58	18.96	19.27	14.23	19.02	14.12	17.69
MEMB1001606	8.92	10.53	14.14	3.81	11.8	8.28	8.12	2.83	3.68
MEMB1001612	12.87	18.3	27.94	17.58	18.03	14.45	13.69	13.84	21.14
MEMB1001618	47.24	51.05	53.93	31.05	43.7	40.99	38.7	45.72	63.89
MEMB1001619	21.53	28.8	35.45	21.89	15.8	18.53	19.99	9.27	17.16
MEMB1001623	32.27	35.63	43.91	23.19	18.98	32.29	44.61	25.32	56.22
MEMB1001625	8.67	7.58	12.04	6.34	5.5	6.91	4.26	3.83	6.59
MEMB1001630	10.6	14.13	23.29	7.12	9.43	7.75	11.72	5.74	8.87
MEMB1001635	8.76	10.43	14.63	5.84	10.38	7.68	13.65	6.82	10.88
MEMB1001637	32.58	33.36	51.7	33.85	36.88	33.36	45.82	19.82	25.12
MEMB1001641	50.01	60.79	92.28	48.36	49.88	43.85	47.93	38.87	68.74
MEMB1001653	12.57	22.18	30.48	17.69	15.29	13.82	24.06	13.06	30.12
MEMB1001665	3.93	2.53	6.42	3.83	1.95	2.63	2.78	1.9	1
MEMB1001665	19.92	21.89	36.37	13.51	17.12	14.78	11.97	10.88	17.21
MEMB1001667	16.89	20.47	51.28	17.31	18.04	14.47	10.43	9.14	14.45
MEMB1001668	4.06	3.82	13.97	2.65	3.74	1.89	2.27	0.07	2.33
MEMB1001669	4.2	4.89	17.85	5.72	6.1	3.19	6.26	1.74	4.48
MEMB1001670	48.37	46.73	48.84	19.58	78.84	42.13	34.32	15.02	19.72
MEMB1001673	67.96	61.75	141.78	71.28	44.3	44.21	62.82	58.55	68.02
MEMB1001675	17.04	14.33	30.97	18.1	10.45	10.58	21.47	8.1	12.88
MEMB1001679	8.24	17.91	21.95	5.19	4.98	3.87	10.86	6.26	8.43
MEMB1001684	10.61	21.58	17.28	4.97	8.8	7.91	15.07	5.39	2.51
MEMB1001685	5.57	8.99	14.97	3.84	6.75	3.87	5.52	1.85	2.86
MEMB1001695	5.09	9.94	17.45	5.82	8.3	5.19	6.68	2.16	2.97
MEMB1001703	4.68	6.84	16.43	5.87	6.69	4.05	8.72	2.46	4.94
MEMB1001704	5.61	7.88	21.01	10.94	17.98	9.05	14.4	6.3	9.45
MEMB1001705	7.42	2.91	5.37	4.07	5.68	4.18	6.33	3.38	3.19
MEMB1001707	3.87	5.88	5.44	3.17	4.08	5.01	5.27	1.48	7.88
MEMB1001717	14.88	12.78	30.35	13.57	11.78	8.58	13.5	11.39	12.48
MEMB1001731	39.25	68.03	95.93	39.11	36.32	32.38	48.55	35.06	44.86
MEMB1001734	5.17	6.48	12.05	1.52	7.18	2.49	5.84	1.69	6.44

34 / 157

2506

MEMB1001456	8.69	12.18	21.38	34.84	17.29	10.08	13.2	9.97	12.41
MEMB1001461	19.67	23.1	39.66	20.5	28.42	24.75	17.54	14.4	25.15
MEMB1001463	19.58	21.33	35.61	31.93	33.77	43.28	30.72	20.88	36.42
MEMB1001464	28.01	27.99	31	14.82	14.82	16.04	16.45	13.58	23.87
MEMB1001466	16.16	16.92	27.63	19.85	13.38	15.95	13.7	10.31	15.02
MEMB1001482	17.23	20.81	35.32	19.01	19	19.25	19.83	14.79	21.88
MEMB1001500	41.82	45.64	68.26	37.83	38.48	37.77	46.2	37.01	39.45
MEMB1001505	20.94	22.59	32.35	28.78	22.08	24.15	26.64	17.19	32.69
MEMB1001521	8.53	7.82	10.6	8.16	6.3	6.64	6.41	3.73	7.14
MEMB1001527	2.99	3.88	4.38	4.6	7.15	3.37	3	1.09	4.09
MEMB1001530	3.48	3.01	4.27	1.8	4.75	3.13	7.05	4.01	3.15
MEMB1001531	36.17	50.68	53.68	24.43	15.05	19.88	14.37	19.2	25.31
MEMB1001532	17.53	24.2	32.45	18.39	15.75	13.5	15.22	13.76	17.31
MEMB1001536	12.38	16.91	23.89	12.47	13.36	10.82	9.04	12.27	13.78
MEMB1001538	66.38	84.8	143.72	91.2	78.03	69.03	101.64	71.92	118.47
MEMB1001537	11.79	11.39	16.93	9.29	11.02	10.91	6.73	5.87	10.85
MEMB1001542	14.98	22.23	29.71	13.95	16.64	16.35	12.71	5.92	15.48
MEMB1001543	23.68	28.65	30.69	20.28	24.21	31.58	17.34	19.68	58.07
MEMB1001547	23.94	50.04	39.45	14.74	15.27	12.72	12.08	2.84	4.52
MEMB1001548	3.11	4.4	5.6	1.9	1.94	2.29	3.65	4.19	3.74
MEMB1001551	3.8	7.07	10.09	4.84	4.24	4.26	3.14	1.85	6.18
MEMB1001555	8.19	9.59	15.51	4.72	6.09	5.64	3.45	5.22	5.59
MEMB1001562	17.05	18.48	38.18	19.29	13.38	17.57	18.22	12.15	22.52
MEMB1001564	38.37	50	64.37	49.41	38.27	48.75	40.56	25.48	48.57
MEMB1001565	18.11	21.08	35.2	23.9	18.29	10.69	18.09	9.8	15.5
MEMB1001566	18.83	22.88	28.98	20.6	18.61	17.83	18.18	11.82	14.18
MEMB1001572	3.68	5.99	8.15	7.26	5.87	2.52	10.81	4.11	5.2
MEMB1001585	31.65	42.11	57.35	32.5	28.22	32.05	31.45	26.22	33.57
MEMB1001585	19.82	27.08	60.2	25.12	17.34	21.21	21.42	13.56	26.31
MEMB1001588	3.6	1.4	8.8	2.53	0.56	1.61	2.28	0.38	3.78
MEMB1001595	18.38	11.37	25.05	8.89	10.88	9.92	9.72	7.65	13.65
MEMB1001595	50.23	70.48	94.67	85.43	78.99	83.08	60.67	47.9	67.78
MEMB1001599	12.08	19.55	22.95	3.76	11.11	7.77	4.14	3.06	5.8
MEMB1001603	15.68	23.5	36.58	18.96	19.27	14.23	19.02	14.12	17.69
MEMB1001606	8.92	10.53	14.14	3.81	11.8	8.28	8.12	2.83	3.68
MEMB1001612	12.87	18.3	27.94	17.58	18.03	14.45	13.69	13.84	21.14
MEMB1001618	47.24	51.05	53.93	31.05	43.7	40.99	38.7	45.72	63.89
MEMB1001619	21.53	28.8	35.45	21.89	15.8	18.53	19.99	9.27	17.16
MEMB1001623	32.27	35.63	43.91	23.19	18.98	32.29	44.61	25.32	56.22
MEMB1001625	8.67	7.58	12.04	6.34	5.5	6.91	4.26	3.83	6.59
MEMB1001630	10.6	14.13	23.29	7.12	9.43	7.75	11.72	5.74	8.87
MEMB1001635	8.76	10.43	14.63	5.84	10.38	7.68	13.65	6.82	10.88
MEMB1001637	32.58	33.36	51.7	33.85	36.88	33.36	45.82	19.82	25.12
MEMB1001641	50.01	60.79	92.28	48.36	49.88	43.85	47.93	38.87	68.74
MEMB1001653	12.57	22.18	30.48	17.69	15.29	13.82	24.06	13.06	30.12
MEMB1001665	3.93	2.53	6.42	3.83	1.95	2.63	2.78	1.9	1
MEMB1001665	19.92	21.89	36.37	13.51	17.12	14.78	11.97	10.88	17.21
MEMB1001667	16.89	20.47	51.28	17.31	18.04	14.47	10.43	9.14	14.45
MEMB1001668	4.06	3.82	13.97	2.65	3.74	1.89	2.27	0.07	2.33
MEMB1001669	4.2	4.89	17.85	5.72	6.1	3.19	6.26	1.74	4.48
MEMB1001670	48.37	46.73	48.84	19.58	78.84	42.13	34.32	15.02	19.72
MEMB1001673	67.96	61.75	141.78	71.28	4				

2507

2508

HENBB1001735	41.8	74.33	131.1	48.7	40.13	60.42	63.81	28.45	83.52
HENBB1001736	7.77	13.39	28.59	8.57	9.16	5.35	8.09	5.45	9.42
HENBB1001747	2.69	8.82	13.53	4.41	6.88	3.98	4.86	1.06	3.48
HENBB1001749	10.28	24.28	26.11	5.15	8.2	10.2	5.99	4.27	10.01
HENBB1001753	10.3	11.05	20.9	9.25	9.8	9.74	6.82	6.38	10.12
HENBB1001756	6.53	8.08	12.88	3.8	3.99	4.72	5	3.97	3.65
HENBB1001767	7.8	24.09	26.61	6.4	12.83	5.98	11.18	14.35	9.41
HENBB1001760	14.27	19.82	42.08	14.61	14.45	11.21	17.24	9.18	15.44
HENBB1001762	0.8	3.04	8.27	0.57	4.82	0.71	2.56	0.81	3.52
HENBB1001780	13.76	23.09	49.83	16.38	14.53	12.62	15.44	6.8	13.71
HENBB1001785	6.49	12.38	24.07	15.82	12.17	15.29	9.31	2.84	7.32
HENBB1001788	25.43	19.69	45.01	19	16.71	23.01	19.54	20.38	28.43
HENBB1001793	5.14	15.25	7.48	1.07	12.29	8	16.96	6.27	6.44
HENBB1001797	14.78	27.69	37.34	12.45	18.08	22.63	22.37	14.75	11.18
HENBB1001802	13.85	8.42	12.11	8	9.1	3.62	4.89	7.62	4.73
HENBB1001812	19.94	26.82	51.97	30.77	23.99	24.59	25.74	15.27	19.71
HENBB1001815	11.61	15.53	37.64	17.99	17.45	14.83	20.39	13.64	19.43
HENBB1001816	17.77	23.19	75.37	26.22	24.8	18.08	30.17	15.34	21.15
HENBB1001831	8.13	9.07	23.11	9.34	10.75	10.45	10.25	3.73	10.62
HENBB1001834	14.4	18	35.74	22.13	11.64	19.05	15.65	10.35	12.6
HENBB1001838	20.28	29.5	43.48	42.55	36.92	34.82	31.2	16.99	16.41
HENBB1001839	14.35	16.68	38.4	17.83	18.75	14.74	12.83	10.7	10.84
HENBB1001841	25.77	32.45	61.17	33.9	28.13	25.07	27.91	22.04	19.78
HENBB1001844	15.48	34.84	44.55	31	24.08	23.04	21.12	16.3	25.7
HENBB1001847	17.89	27.69	60.83	26.2	23.47	23.52	21.19	9.97	20.24
HENBB1001848	4.85	5.43	11.29	3.58	4.51	3.65	6.07	3.08	5.61
HENBB1001850	38.14	52.12	78.88	56.18	68.55	53.58	61.43	34.94	43.48
HENBB1001859	30.81	27.05	60.38	58.19	35.48	57.28	19.8	14.26	20.24
HENBB1001863	3.48	3.88	8.32	5.64	4.36	4.8	7.82	2.45	4.74
HENBB1001867	14.5	16.62	28.09	12.79	10.98	12.92	14.13	8.77	15.8
HENBB1001868	5.84	8.18	12.72	7.12	6.24	5.61	8.09	8.94	8.48
HENBB1001869	54.88	64.59	94.37	86.31	67.88	82.85	80.48	51.3	89.82
HENBB1001872	6.07	11.88	13.6	18.68	9.48	7.21	18.78	6.55	10.02
HENBB1001874	3	3.3	5.61	4.59	12.05	11.19	3.13	1.36	5.74
HENBB1001875	12.99	18.2	23.61	26.62	29.44	24.9	34.89	17.55	28.63
HENBB1001880	18.04	17.79	23.89	31.27	17.83	20.15	14.84	10.37	14.08
HENBB1001899	34.29	46.59	83.32	39.63	32.89	31.82	35.58	26.2	35.72
HENBB1001903	8.18	10.41	29.09	8.47	10.89	12.17	9.48	4.66	7.91
HENBB1001905	28.16	29.19	50.5	33.48	32.78	32.37	33.69	21.8	33.71
HENBB1001905	4.65	5.05	9.2	6.27	6.88	4.65	7.25	6.26	9.65
HENBB1001908	2.42	5.7	5.4	5.21	4.63	6.7	4.19	0.96	5.33
HENBB1001910	5.2	10.82	10.84	8.51	10.51	9.98	6.79	2.78	7.39
HENBB1001911	29.98	51.31	56.74	43.98	49.18	34.13	49.74	26.26	62.6
HENBB1001915	21.14	31.44	37.82	20.71	15.99	14.9	18.9	21.98	29.35
HENBB1001921	7.15	10.68	16.34	7.38	7.15	5.68	5.97	5.19	7.01
HENBB1001922	17.41	22.59	26.93	13.84	10.93	18.21	23.02	11.95	17.79
HENBB1001925	13.87	12.3	28.77	11.6	11.78	10.4	15.51	12.85	15.94
HENBB1001930	40.03	69.19	73.92	44.03	41.91	43.37	40.7	31.14	54.53
HENBB1001944	1.94	5.58	8.4	6.68	4.5	5.15	2.97	2.18	4.4
HENBB1001845	11.9	14.15	28.25	12.81	15.86	20.85	15.45	10.36	15.75
HENBB1001847	12.74	2.15	16.64	14.03	14.01	12.25	15.71	4.99	12.24
HENBB1001850	17.57	25.59	40.32	25.51	11.85	16.5	13.59	12.65	18.84
HENBB1001852	26.14	33.12	47.22	23.32	21.94	24.22	34.78	37.99	49.1
HENBB1001853	17.2	26.64	33.2	20.18	17.54	17.43	18.02	10.46	24.03
HENBB1001857	5.93	5.68	12.29	2.87	5.82	5.6	4.01	5.73	6.82
HENBB1001859	4.28	3.68	11.59	1.82	2.82	5.97	1.57	2.64	3.72
HENBB1001862	11.92	43.87	27.13	12.9	32.94	14.36	12.1	15.99	21
HENBB1001867	34.68	44.03	60.34	44.77	38.71	38.73	44.26	27.2	37.33
HENBB1001873	28.25	35.18	36.62	28.87	29.33	21.67	37.28	18.87	23.03
HENBB1001874	70.1	95.85	135.82	108.46	60.16	80.87	54.72	81.03	88.68
HENBB1001881	4.69	4.08	8.04	3.84	5.22	5.54	5.15	3.31	5.57
HENBB1001887	10.65	12.27	28.1	12.68	12.32	11.95	11.85	6.64	23.87
HENBB1001888	9.33	7.22	29.4	8.24	11.34	8.88	13.31	6.58	9.69
HENBB1001890	26.2	19.68	27.78	43.3	26.68	36.79	27.71	13.13	24.3

35 / 157

2509

2510

MEMB1001986	27.04	42.61	48.67	17.55	24.82	18.56	28.17	23.4	25.38	*	-
MEMB1001987	14.71	21.01	29.3	15.9	17.15	12.6	18.88	11.27	17.5		
MEMB1001989	15.2	22.48	20.61	18.58	17.9	14.38	19.76	8.32	11.63		
MEMB1002002	6.41	8.68	14.02	3.88	4.69	7.48	2.32	2.31	9.35		
MEMB1002005	13.93	14.9	24.54	14.07	12.64	13.23	24.2	15.6	24.4		
MEMB1002009	13.81	20.37	28.6	12.9	13.24	9.2	14.63	11.53	16.62		
MEMB1002013	35.52	43.43	61.55	22.34	25.62	25.39	38.97	23.31	38.2	*	-
MEMB1002015	17.64	18.37	28.67	10.62	13.94	14.86	13.44	9	13.73		
MEMB1002024	5.8	10.6	13.79	2.3	5.04	3.13	2.77	1.85	3.6	*	-
MEMB1002035	52.65	85.52	32.75	21.71	70.63	38.58	40.44	20.7	17.84		
MEMB1002039	4.96	6.73	12.89	4.98	10.04	5.87	7.91	2.61	3.57		
MEMB1002041	26.99	42.84	59.33	33.42	31.41	30.59	35.6	25.94	38.68		
MEMB1002042	7.32	8.11	15.44	8.43	6.37	4.7	6.95	4.45	6.64		
MEMB1002043	15.38	16.33	33.37	9.15	15.15	12.75	12.63	10.91	12.91		
MEMB1002044	8.28	7.19	15.99	4.83	6.03	3.49	8.95	3.61	5.78		
MEMB1002045	13.55	14.05	38.81	21.11	17.88	16.3	14.27	7.83	17.45		
MEMB1002049	20.72	21.4	45.82	13.98	20.22	15.6	21.11	13.82	17		
MEMB1002050	12.83	16.25	37.04	11.52	18.48	11.6	18.41	9.54	14.5		
MEMB1002051	28.94	18.92	52.44	23.53	25.05	15.13	33.91	14.41	28.65		
MEMB1002068	64.68	61.13	134.22	59.82	39.3	49.08	48.07	48.15	30.52		
MEMB1002069	20.88	27.08	35.85	26.67	17.79	18.95	23.6	16.88	23.73		
MEMB1002075	15.68	17.27	22.02	11.81	12.44	12.63	5.87	8.58	11.32	*	-
MEMB1002079	27.48	41.14	33.47	23.15	21.35	9.24	13.25	11.83	12.65	**	-
MEMB1002080	27.32	43.03	71.5	48.33	38.81	19.92	37.68	22.98	28.59		
MEMB1002082	11.54	12.93	24.88	5.24	21.72	15.15	15.69	4.71	14.1		
MEMB1002084	17.19	20.49	48.44	15.57	24.58	30.61	24.15	14.34	22.35		
MEMB1002088	4.85	8.19	14.68	4.12	8	4.95	6.55	2.38	8.06		
MEMB1002092	144.34	118.11	135.03	45.77	56.49	60.48	15.5	25.51	16.94	**	-
MEMB1002094	15.12	19.26	35.12	17.78	15.07	8.68	14.96	8.23	22.31		
MEMB1002103	5.48	7.53	11.18	9.4	4.51	4.6	8.52	1.47	4.97		
MEMB1002109	59.58	83.48	107.06	62.19	60.92	51.12	65.01	50.9	42		
MEMB1002115	5.46	60.72	14.77	19.13	18.84	5.85	5.92	6.74	5.3		
MEMB1002120	6.4	5.75	12.8	2.74	5.51	2.84	3.13	1.97	3.49		
MEMB1002121	19.45	25.43	55.52	28.58	24.08	19.83	25.58	15.89	20.17		
MEMB1002134	10.91	24.24	48.39	24.03	23.78	24.09	19.26	8.44	14.1		
MEMB1002136	18.33	18.78	45.57	23.81	13.25	20.44	11.9	15.85	24.31		
MEMB1002138	58.09	51.59	110.11	35.51	42.15	39.79	27.69	28.58	25.22		
MEMB1002139	8.36	15.47	10.43	4.01	4.89	4.74	7.89	6.83	4.81	*	-
MEMB1002141	9.19	16.96	13.1	1.9	10.98	7	5.85	7.53	4.35	*	-
MEMB1002142	8.38	8.37	8.62	2.74	2.82	2.25	4.04	2.02	5.71	**	-
MEMB1002145	18.49	18.24	32.26	9.14	12.19	10.34	12.54	5.18	7.82		
MEMB1002162	24.88	22.9	59.51	8.84	21.68	20.2	34.39	10.44	7.82		
MEMB1002162	46.61	80.75	107.44	68.24	81.5	44.12	82.89	47.69	70.78		
MEMB1002173	3.55	25.4	7.84	5.45	25.88	14.78	4.91	5.53	5.89		
MEMB1002188	5.91	5.38	11.91	4.49	4.74	3.73	7.59	4.5	2.15		
MEMB1002190	8.05	5.19	7.81	3.82	3.51	3.24	3.4	1.73	2.03	*	-
MEMB1002193	13.5	31.42	40.95	12.51	15.71	29.81	30.17	23.34	31.48	*	-
MEMB1002217	11.78	15.58	34.59	12.72	11.26	16.37	17.63	11.07	12.8		
MEMB1002218	25.74	52.63	73.43	32.82	28.59	24.52	35.6	20.84	26.88		
MEMB1002228	188.24	84.59	214.25	80.82	78.38	108.65	188.71	122.42	65.1		
MEMB1002232	10.76	24.39	31.98	19.41	21.33	17.37	22.17	5.39	14.18		
MEMB1002245	29.32	39.47	46.03	33.89	25.42	30.62	28.25	20.38	28.6		
MEMB1002247	23.04	30.4	36.26	21.55	23.11	19.18	30.01	21.82	18.27		
MEMB1002248	8.45	5.4	12.9	9.1	8.18	7.68	2.82	2.96	4.18		
MEMB1002254	13.65	8.54	22.22	9.35	11.14	10.19	5.38	9.93	4.18		
MEMB1002255	24	39.55	56.2	39.15	37.05	37.31	39.64	23.55	35.88		
MEMB1002256	19.55	37.2	69.12	27.4	30.03	37.09	27.79	17.42	23.42		
MEMB1002271	8.4	14.75	33.73	16.63	15.77	17.75	17.9	8.81	11.82		
MEMB1002280	0	1.32	3.2	1.59	2.57	0.19	2.25	0	1.09		
MEMB1002286	13.25	17.07	48.86	29.84	17.48	26.16	17	14.47	20.56		
MEMB1002300	7.2	17.98	22.84	22.43	7.83	12.63	8.34	8.09	9.04		
MEMB1002302	8.19	9.95	13.55	11.87	6.29	9.78	7.21	2.24	3.19	*	-
MEMB1002306	4.48	4.94	10.04	8.42	4.47	9.84	5.18	3.92	6.24		
MEMB1002316	9.41	18.69	11	13.6	12.17	16.43	7.32	2.42	8.98		

38 / 157

【0860】

【表389】

2511

2512

MEMB1002325	6.79	10.97	11.32	21.02	12.83	6.78	6.12	3.85	6.53
MEMB1002327	29.86	39.37	46.64	28.76	34.27	35.49	23.78	23.05	28.43
MEMB1002329	12.21	17.09	28.97	18.73	8.27	19.71	9.42	11.59	11.1
MEMB1002340	3.57	5.21	5.98	7.77	3.22	9.25	5.09	2.16	1.79
MEMB1002342	6.21	7.8	10.28	8	3.29	3.42	3.33	3.85	5.16
MEMB1002355	32.09	38.53	65.28	39.45	32.86	34.65	33.44	21.57	27.85
MEMB1002356	3.96	3.89	6.45	3.93	4.64	4.39	3.95	0.81	2.74
MEMB1002364	3.43	4.07	8.13	6.97	4.3	5.26	4.27	1.48	3.01
MEMB1002366	7.13	10.16	14.56	9.95	9.83	9.69	9.48	8.8	10.03
MEMB1002371	17.53	29.76	21.42	21.87	21.71	16.54	9.41	9.09	4.87
MEMB1002381	7.92	12.33	16.15	8.95	3.96	8.62	4.76	5.05	4.4
MEMB1002382	135.68	179.98	228.28	288.57	191.47	226.83	107.86	116.89	120.62
MEMB1002387	23.79	23.11	43.65	20.25	12.08	15.04	3.85	6.21	4.42
MEMB1002409	13.31	12.51	22.34	8.77	11.21	10.89	4.44	4.37	4.87
MEMB1002413	3.49	3.12	9.08	4.81	2.78	4	1.1	1.88	1.88
MEMB1002415	2.64	3.61	7.39	2.66	2.05	3.42	0.88	1.28	2.41
MEMB1002424	6.07	9.92	9.97	6.37	9	12.89	8.72	4.79	6.15
MEMB1002425	4.48	7.81	11.5	5.69	9.56	5.52	4.13	1.79	5.11
MEMB1002427	29.25	30.49	50.22	43.33	23.55	33.03	24.65	22.4	31.01
MEMB1002442	6.2	17.54	17.42	9.79	7.28	9.01	4.38	12.11	11.81
MEMB1002447	110.71	207.61	155.4	83.72	84.2	90.9	28.65	37.71	31.85
MEMB1002453	5.36	5.88	9.95	2.72	2.55	3.68	2.41	1.81	3.85
MEMB1002467	16.89	18.27	19.17	8.78	17.09	11.88	7.07	5.54	6.6
MEMB1002468	23.11	31.15	52.28	31.43	27.93	29.52	25.75	14.57	26.97
MEMB1002483	24.77	42.25	57.37	32.51	37.48	24.85	28.18	20.41	30.3
MEMB1002486	3.27	5.17	9.01	3.69	4.27	4.31	3.51	0.9	2.26
MEMB1002477	4.25	7.51	5.14	7.08	3.76	4.24	1.88	3.33	2.58
MEMB1002479	20.58	32.93	33.42	17.09	21.13	16.76	11.95	15.32	21.36
MEMB1002489	48.69	62.6	67.58	34.04	44.48	57.32	35.3	38.02	39.44
MEMB1002492	11.53	17.66	26.19	13.16	10.94	12.55	11.2	5.85	16.28
MEMB1002495	5.18	3.21	8.5	2.92	4.34	3.1	3.82	2.25	3.88
MEMB1002502	34.71	31.95	38.03	48.31	31.48	35.49	28.54	23.82	25.15
MEMB1002509	11.07	15.65	26.84	12.73	16.51	11.29	10.77	7	12.74
MEMB1002510	28.51	45.37	36.84	31.94	38.1	26.25	23.53	21.31	17.08
MEMB1002520	14.34	15.82	18.28	10.3	7.17	4.37	1.27	2.73	3.86
MEMB1002522	27.23	55.84	54.45	41.8	39.8	28.31	15.32	22.78	36.38
MEMB1002527	2.58	2.91	5.88	1.88	2.03	3.34	1.71	1.04	2.87
MEMB1002530	5.18	6.33	11.67	5.26	3.42	5.09	4.28	2.92	7.5
MEMB1002531	26.73	31.41	45.82	22.42	21.99	29.07	24.79	13.7	18.84
MEMB1002534	27.18	48.28	57.65	25.13	29.78	23.09	34.17	26.17	35.08
MEMB1002536	13.04	20.55	23.13	9.22	13.07	13	15.33	12.24	18.4
MEMB1002544	23	32.2	43.29	21.03	20.23	27.52	19.17	15.09	14.85
MEMB1002545	3.59	4.33	9.7	5.81	3	3.66	1.12	0.98	13.79
MEMB1002560	7.33	13.21	13.27	3.83	8.6	4.73	2.81	3.81	2.61
MEMB1002556	19.38	24.38	22.91	7.86	11.05	11.87	7.3	3.24	5.41
MEMB1002571	7.32	4.72	8.5	4.01	3.71	2.33	1.87	1.16	2.04
MEMB1002579	3.21	4.8	10.5	2.3	2.84	2.3	2.51	1.04	2.73
MEMB1002582	9.95	12.82	19.84	6.38	8.65	6.74	3.38	2.01	3.49
MEMB1002584	4.7	5.64	13.38	4.24	5.58	6.79	3.53	1.48	2.34
MEMB1002587	8.59	15.17	29.93	11.81	13.87	11.06	10.85	4.49	11.04
MEMB1002590	27.91	31.34	78.12	45.05	22.1	24.36	21.48	22.81	39.4
MEMB1002596	9.4	9.38	12.16	12.87	8.82	9.84	3.39	4.59	5.2
MEMB1002600	7.35	8.89	8.83	4.2	3.37	4.13	3.83	2.53	0.85
MEMB1002601	13.59	25.38	25.69	13.58	12.88	10.65	15.53	14.47	12.11
MEMB1002603	4.01	4.35	8.49	3.89	3.23	2.33	3.4	1.13	1.07
MEMB1002607	5.58	8.76	12.05	6.25	8.3	3.46	5.04	2.83	4.29
MEMB1002610	6.17	9.3	18.74	7.02	8.85	5.18	5.21	2.01	3.82
MEMB1002613	5.1	6.85	16.57	4.47	6.62	3.43	4.07	1.86	3.63
MEMB1002614	188.18	118.26	257.79	180.19	127.61	115.27	80.95	70.77	100.61
MEMB1002615	3.78	4.74	6.46	3.53	4.89	3.11	2.31	1.58	0.85
MEMB1002617	15.34	18.43	20.69	14.08	10.48	7.36	18.03	9.38	12.23
MEMB1002623	43.48	62.43	70.11	89.32	59.95	84.13	48.57	40.8	51.16
MEMB1002624	20.45	39.7	56.1	29.45	24.85	24.11	26.5	22.1	22.85
MEMB1002631	12.8	24.72	22.26	5.67	13.71	7.87	12.48	14.8	9.64

37 / 157

【0861】

【表390】

2513

2514

MEMB1002485	3.81	5	11.37	1.21	3.62	1.08	2	1.05	2.82										
MEMB1002444	8.38	7.38	19.6	8.87	8.62	5.91	7.04	3.57	7.83										
MEMB1002454	44.15	47.27	88.13	55.08	50.78	51.15	27.74	20.25	41.98										
MEMB1002481	16.29	22.35	33.51	18.43	18.3	14.62	15.42	10.61	19.12										
MEMB1002463	12.29	20.14	24.54	5.77	16.48	12.72	3.86	7.87	3.2										
MEMB1002464	26.04	27.32	31.87	12.71	14.38	12.28	15.87	10.5	16										
MEMB1002477	2.85	4.87	7.15	1.19	3.8	1.22	2.58	0.54	2.04										
MEMB1002483	408.12	630.85	681.04	161.92	536.41	322.12	493.81	190.86	318.34										
MEMB1002484	5.89	14.15	21.74	5.84	5.87	5.5	2.78	0.53	2.33										
MEMB1002488	2.08	2.21	13.73	0.94	2.87	1.98	2.53	0.03	2.4										
MEMB1002492	6.27	8.28	18.48	7.58	4.81	4.6	4.23	1.22	6.95										
MEMB1002493	37.13	59.89	60.13	8.18	8.08	32.77	12.89	24.23	25.51										
MEMB1002497	12.19	19.73	29.39	11.74	11.32	12.95	13.01	7.88	10.37										
MEMB1002499	12.29	7.07	29.87	3.99	12.05	3.69	6.19	2.34	1.91										
MEMB1002702	4.21	4.82	9.81	4.02	2.81	2.59	2.32	3.45	2.32										
MEMB1002705	42.01	58.09	92.41	42.52	47.93	35.39	82.82	52.43	59.52										
MEMB1002712	82.07	113.22	149.82	77.86	93.78	72.72	96.78	72.75	88.71										
MEMB1000028	2.68	4.75	13.62	4.72	4.88	2.52	2.74	1.07	1.92										
MEMB1000031	7.53	10.04	0	0.25	16.99	0.79	7.55	3.43	9.54										
MEMB1000034	9.42	10.78	32.44	15.75	18.03	17.48	8.88	5.24	7.94										
MEMB1000039	1.72	2	2.81	3.07	3.78	1.79	0.78	0	1.77										
MEMB1000044	5.65	7.08	18.45	11.11	7.14	7.03	4.83	2.88	3.82										
MEMB1000049	10.51	11.67	19.12	13.18	11.8	12.05	6.47	7.76	7.58										
MEMB1000055	3.33	5.05	8.68	2.81	3.29	3.45	1.85	0.92	1.07										
MEMB1000059	3.15	5.89	14.29	4.37	6.53	5.77	1.28	1.07	2.45										
MEMB1000091	59.8	78.47	131.41	171.82	125.28	143.68	43.76	27.21	45.98										
LIVER1000034	4.13	4.93	5.75	3.08	2.78	4.02	2.18	1.07	1.38										
LIVER1000038	10.35	13.8	22.11	11.7	10.53	14.7	11.26	5.83	10.54										
LIVER1000111	59.6	50.92	129.92	124.27	66.83	94.45	48.23	37.88	48.08										
LIVER1000222	2.42	3.87	7.13	5.18	2.89	5.96	3.89	0.85	2.2										
LIVER1000225	11.19	21.41	14.85	14.92	17.52	18.14	8.19	3.83	7.48										
LIVER1000330	5.78	4.25	9.83	9.98	5.67	5.82	4.77	2.89	3.83										
LIVER1000445	4.77	5.88	5.47	6.22	7.71	6.61	4.81	3.96	4.97										
LIVER1000446	23.39	30.93	40.35	28.87	14.67	17.18	8.83	19.03	18.73										
LIVER1000772	9.51	20.72	24.34	17.51	14.57	15.78	8.13	9.06	13.17										
LIVER1000777	7.51	7.91	13.85	13.48	8.07	9.05	5.23	4.4	6.07										
LIVER1000880	5.07	5.78	10.98	6.47	4.34	6.39	3.86	2.15	4.58										
LIVER1000885	11.03	13.61	23.45	8.67	23.05	12.3	11.48	8.1	16.78										
LIVER1000992	19.59	15.75	23.53	19.18	22.81	13.71	8.85	6.69	7.35										
LIVER1000995	61.53	68.53	98.85	69.98	49.48	69.44	74.79	43.47	64.56										
LIVER1000997	8.95	15.6	14.45	10.75	13.82	9.14	5.8	2.85	3.48										
LIVER1000998	41.49	40.97	103.47	50.68	24.2	48.28	26.16	32.27	27.07										
LIVER1001000	3.55	4.13	7.44	3.5	3.68	3.78	2.58	3.54	3.3										
LIVER1001001	44.76	62.72	56.32	38.92	31.17	34.9	40.25	35.88	28.75										
LIVER1001006	3.83	4.77	8.45	2.61	3.61	4.62	2.14	3.23	3.22										
LIVER1001008	6.19	8.29	14.94	4.01	4	8.54	5.47	8.84	5.37										
LIVER1001115	1.31	3.43	5.11	1.07	1.91	1.17	0.78	1.21	1.27										
LIVER1001120	5.87	8	12.35	7.59	6.45	6.85	5.23	2.59	6.22										
LIVER1001138	5.5	7.3	19.82	7.01	10.38	6.37	5.23	4.12	9.23										
LIVER1001148	4.13	5.08	8.45	8.16	4.26	4.91	1.55	2.41	3.37										
LIVER1001148	10.81	16.45	14.93	9.88	11.05	13.74	8.11	8.29	5.5										
LIVER1001157	4.51	3.83	8.67	3.38	5.06	4.59	1.48	2.24	4.29										
LIVER1001181	85.1	101.44	127.49	155.4	95.91	123.47	53.19	53.6	81.33										
LIVER1001187	9.34	7.8	16.08	6.28	7.38	6.18	4.07	3.11	4.06										
LIVER1001174	14.38	23.84	29.19	10.08	10.64	10.6	4.27	1.62	4.28										
LIVER1001185	15.55	11.69	24.81	3.88	12.76	7.24	8.4	3.76	4.58										
LIVER1001187	40.28	72.32	62.82	52.19	53.14	48.8	30.48	15.14	10.77										
LIVER1001190	13.01	13.24	16.57	5.85	5.19	5.07	2.85	4.77	5.27										
LIVER1001192	3.52	6.68	4.51	4.92	4.75	3.34	0.75	1.7	3.55										
MEMB1000009	5.18	9.12	9.31	4.2	4.18	3.51	1.94	1.4	2.13										
MEMB1000015	5.78	4.65	11.74	5.4	3.83	5.38	2.48	1.85	3.15										
MEMB1000019	8.71	5.88	17.78	9.05	5.32	6.22	3.67	4.83	8.34										
MEMB1000020	101.37	73.82	112.58	134.32	82.42	108.51	82.61	44.53	81.2										
MEMB1000024	19.83	38.9	35.17	7.48	24.94	15.61	12.77	15.51	8.89										

38 / 157

2515

2516

RAMBA 1000015	3.49	6.73	9.73	4.76	8.15	5.09	1.99	1.9	3.02
RAMBA 1000043	8.59	8.02	15.22	6.68	8.56	4.78	2.11	1.9	8.7
RAMBA 1000045	6.03	8.88	17.21	11.41	6.71	9.34	5.16	4.7	6.29
RAMBA 1000046	41.47	58.57	56.97	30.34	27.99	31.58	37.45	30.61	48.64
RAMBA 1000055	10.6	18.69	12.32	4.6	6.26	5.68	3.62	4.4	2.53
RAMBA 1000057	3.76	2.41	12.19	2.23	1.8	1.43	1.21	1.53	2.81
RAMBA 1000101	10.21	8.2	18.54	5.99	7.11	5.68	5.98	4.17	6.29
RAMBA 1000103	143.75	270.15	259.45	48.26	67.48	38.63	33.38	39.16	49.31
RAMBA 1000018	3.34	5.16	10.33	3.47	4.79	2.05	3	0.87	2.68
RAMBA 1000035	8.66	4.9	7.86	4.66	4.18	4.45	2.38	1.98	9.84
RAMBA 1000032	1.86	4.62	8.88	3.95	2.75	3.09	2.27	0.07	1.18
RAMBA 1000036	3.52	3.48	7.14	2.58	2.53	2.5	0.67	1.83	1.25
RAMBA 1000037	4.98	8.73	8.88	2.64	2.02	1.49	1.93	0.71	1.64
RAMBA 1000102	16.31	20.65	19.78	15.97	14.11	11.62	28.48	21.84	17.28
RAMBA 1000103	24.25	30.87	39.49	18.45	29.7	17.72	21.24	13.56	7.61
RAMBA 1000105	8.95	10.76	22.4	12.82	11.13	10.63	6.32	4.47	6.45
RAMBA 1000117	2.83	6.08	12.12	2.65	4.11	2.11	5.97	1.27	1.18
RAMBA 1000118	8.06	8.77	19.68	14.56	12.82	9.23	5.17	8.7	14.06
RAMBA 1000129	4	2.76	6.79	3.48	3.6	1.79	1.7	2.01	2.69
RAMBA 1000132	5.05	7.04	8.39	3.71	2.81	3.3	8.78	9.22	3.38
RAMBA 1000134	7.25	6.17	12.23	3.97	5.48	2.89	5.9	4.84	5.79
RAMBA 1000139	8.61	13.92	27.94	9.57	7.21	7.44	10.8	7.32	10.69
RAMBA 1000141	10.59	12.38	22.85	16.48	15.91	12.48	8.35	6.81	8.58
RAMBA 1000143	47.45	88.55	73.36	60.42	67.69	52.39	70.22	28.61	57.3
RAMBA 1000150	7.57	5.6	14.19	2.79	8.22	2.73	5.8	1.58	3.48
RAMBA 1000155	11.41	12.76	22.27	10.11	6.82	6.05	2.76	1.55	4.85
RAMBA 1000163	15.02	19.25	13.92	12.25	12.39	13.68	10.19	9.3	14.14
RAMBA 1000171	7.6	10.37	13.24	2.98	2.32	1.89	2.54	1.28	1.33
RAMBA 1000173	6.75	5.29	9.57	2.32	2.23	1.04	2.07	1.37	0.89
RAMBA 1000175	3.48	6.59	11.02	3.52	2.3	2.43	3.53	1.51	2.66
RAMBA 1000183	25.33	42.64	44.08	23.65	35.89	24.05	15.46	12.47	14.16
RAMBA 1000191	5.92	8.95	18.7	5.23	6.43	8.14	3.97	2.29	5.68
RAMBA 1000192	11.99	14.66	36.16	26.22	18.77	12.98	9.82	7.53	11.64
RAMBA 1000193	1.19	3.37	6.86	4.96	4.82	4.54	1.06	1.39	2.14
RAMBA 1000198	47.48	58.45	45.4	32.82	40.48	48.35	28.71	22.96	42.97
RAMBA 1000204	11.23	12.69	25.92	6.96	11.44	9.78	5.53	5.78	6.1
RAMBA 1000207	12.4	7.47	19.55	6.28	5.69	4.84	6.03	8.39	7.97
RAMBA 1000214	8.42	8.76	13.08	9.58	9.32	7.58	7.33	4.95	7.33
RAMBA 1000221	22.52	15.12	35.89	5.02	6.29	8.82	3.88	3.87	3.09
RAMBA 1000221	10.4	16.84	24.93	5.03	11.07	6.49	6.69	3.21	1.94
RAMBA 1000226	13.64	11.97	26.4	5.26	12.91	9.41	26	12.92	3.5
RAMBA 1000227	14.38	17.15	40.21	28.86	16.67	19.14	13.63	7.29	12.48
RAMBA 1000230	3.53	5.09	7.34	3.64	2.92	3.1	1.76	1.24	2.65
RAMBA 1000241	2.76	3.58	6.29	2.63	1.66	1.65	1.27	0.15	1.78
RAMBA 1000245	6.45	7.89	22.51	9.95	8.51	9.68	6.76	3.1	4.3
RAMBA 1000246	21.28	26.61	55.84	30.22	20.66	19.73	24.19	11.24	13.69
RAMBA 1000247	5.31	4.84	18.73	2.94	4.44	3.42	3.87	1.53	2.1
RAMBA 1000254	2.32	3.23	8.55	2.43	2.71	1.01	0.9	0.97	0.54
RAMBA 1000257	2.97	3.77	9.11	1.37	3.58	0.82	2.27	0.96	1.23
RAMBA 1000262	45.74	36.4	0	0	39	0	34.58	23.42	42.95
RAMBA 1000264	9.24	12.72	35.1	26.8	15.94	23	7.31	7.08	9.28
RAMBA 1000266	0.66	1	3.04	2.93	2.61	0.59	0.15	0	1
RAMBA 1000270	13.12	14.9	18.21	14.67	11.48	11.77	8.4	8.61	6.99
RAMBA 1000271	3.68	3.72	7.35	4.08	3.89	3.79	1.47	8.83	1.69
RAMBA 1000277	41.4	61.52	103.68	30.74	33.8	53.69	17.12	13.62	16.77
RAMBA 1000278	11.35	16.91	37.8	22.57	16.83	18.11	10.35	6.71	12.64
RAMBA 1000279	3.26	3.91	6.49	3.15	3.6	2.66	2.52	0.56	1.77
RAMBA 1000283	28.38	38.95	49.97	68.77	34.21	54.3	27.86	21.79	21.63
RAMBA 1000284	5.55	6.62	7.77	4.76	4.06	5.69	4.18	2.05	4.21
RAMBA 1000287	18.44	27.31	36.41	15.61	4.83	9.39	4.34	3.99	5.29
RAMBA 1000294	14.59	17.3	29.88	18.53	15.7	20.07	17.88	12.1	15.45
RAMBA 1000296	4.08	4.13	7.44	4.05	8.42	8.4	4.25	3.29	4.19
RAMBA 1000302	3.99	3.48	11.09	6.9	6.88	5.91	4.93	2.1	3.71
RAMBA 1000303	1.77	4.18	5.21	2.14	3.15	3.07	3.83	0.6	1.66

39 / 157

2517

2518

RAMMA1000305	32.01	34.24	58.88	37.52	42.02	39.78	30.9	18.25	28.49
RAMMA1000307	7.32	11.17	15.9	9.52	5.29	9	4.83	4.75	5.32
RAMMA1000309	13.21	12.45	16.61	13.37	8.5	8.68	5.45	4.89	8.28
RAMMA1000312	3.8	4.66	7.48	5.51	4.44	4.58	3.65	2.18	2.53
RAMMA1000313	5.23	3.78	8.22	6.32	8.18	4.37	4.05	1.98	2.38
RAMMA1000321	3.57	2.97	7.78	3.29	3.58	6.01	3.84	1.96	2.65
RAMMA1000325	1.93	3.97	7.5	5.14	3.16	3.87	4.24	1.18	2.51
RAMMA1000332	4.65	7.71	8.76	4.7	8.88	8.63	2.97	1.94	3.31
RAMMA1000340	13.8	21.08	20.08	18.65	20.75	20	20.39	11.88	8.29
RAMMA1000348	6.8	5.86	10.24	5.88	4.39	5.28	1.91	2.51	3.44
RAMMA1000358	48.67	74.66	87.35	63.63	37.37	48.65	35.71	39.77	39.88
RAMMA1000368	9.54	13.7	16.74	8.76	9.05	7.26	6.27	7.75	8.04
RAMMA1000380	4.42	6.35	14.28	6.25	8.74	5.72	3.05	4.95	4.78
RAMMA1000381	62.5	108.93	116.65	101.92	117.16	105.88	87.17	73.39	114.81
RAMMA1000383	4.98	6.13	8.18	4.8	3.58	8.25	3.82	2.21	3.44
RAMMA1000370	5.17	12.45	10.14	6.35	7.18	7.58	3.49	1.29	2.83
RAMMA1000371	2.39	4.39	9.15	3.68	6.58	4.68	2.76	1.36	2.71
RAMMA1000372	17.51	21.43	34.07	12.37	14.77	13.19	11.25	17.12	32.9
RAMMA1000385	2.59	3.13	9.36	4.39	3.21	2.85	3.43	3.75	2.78
RAMMA1000388	15.23	18.38	27.37	8.49	18.09	20.18	11.94	20.32	17.28
RAMMA1000395	2.71	2.65	9.67	2.22	1.18	2.27	1.54	2.51	1.57
RAMMA1000402	17.65	14.83	40.38	21.39	34.95	27.07	11.67	5.88	8.61
RAMMA1000403	4.76	5.81	8.95	2.68	5.93	4.73	3.53	2.03	3.28
RAMMA1000410	8.38	10.39	14.35	8.64	10.52	7.1	6.79	5.15	5.67
RAMMA1000413	13.08	31.41	34.59	25.18	28.63	19.73	20.54	9.59	10.58
RAMMA1000414	28.2	35.16	52.64	41.13	33.28	34.97	22.92	18.5	31.45
RAMMA1000416	93.85	138.41	125.21	48.14	53.85	91.17	35.64	84.76	53.2
RAMMA1000421	12.39	14.43	18.48	7.85	12.15	12.39	5.74	4.49	4.48
RAMMA1000422	142.54	131.07	148.31	147.75	137.43	173.28	122.42	112.28	99.77
RAMMA1000423	5.22	2.78	10.5	5.24	3.98	3.14	2.55	0.29	3.52
RAMMA1000424	16.73	20.92	35.59	20.88	21.28	23.97	15.73	11.76	18.32
RAMMA1000428	9.35	15.84	18.54	7.22	10.71	6.67	4.21	3.75	2.1
RAMMA1000431	1.71	4.18	8.38	1.45	4.13	3.04	2.28	2.12	2.5
RAMMA1000432	63.4	133.96	122.67	9.74	51.75	20.93	35.92	64.94	54.98
RAMMA1000437	7.75	15.03	14.39	12.33	7.78	9.82	5.47	8.03	11.07
RAMMA1000444	7.3	9.64	12.3	7.17	8.74	4.04	4.84	3.53	6.39
RAMMA1000446	72.49	65.48	72.84	87.18	57.65	80.67	67.4	45.57	58.28
RAMMA1000449	18.1	22.35	31.93	13.49	17.76	14	17.72	12.22	18.97
RAMMA1000457	18.4	25.87	17.2	2.75	17.71	11.5	15.1	3.75	15.53
RAMMA1000458	7.78	16.72	16.75	9.89	10.94	4.55	4.37	2.84	7.72
RAMMA1000468	16.03	28.35	43.07	20.2	27.28	30.25	21.32	9.68	14.64
RAMMA1000472	5.65	14.63	17.94	8.59	8.34	7.38	4.92	4.41	8.54
RAMMA1000473	18.92	28.23	28.07	15.83	15.85	17.15	17.93	13.07	20.76
RAMMA1000477	3.29	5.76	7.67	2.77	2.69	3.64	2.99	1.75	1.93
RAMMA1000478	3.8	4.66	7.21	2.84	2.45	2.89	2.43	1.21	0.87
RAMMA1000483	141.62	258.33	267.79	88.09	257.9	180.98	159.43	158.7	175.75
RAMMA1000490	11.74	16.92	30.01	11.37	13.37	11.4	15.88	7.97	17.05
RAMMA1000486	29.48	25.33	53.38	30.03	31.05	22.35	19.18	13.04	15.93
RAMMA1000500	4.21	8.78	19.18	2.1	8.22	6.73	7.01	0.97	5.78
RAMMA1000501	3.3	3.92	7.8	5.4	3.37	3.89	2.1	0.03	4.12
RAMMA1000502	8.55	8.62	11.81	8.78	3.91	8.21	3.85	2.45	1.88
RAMMA1000506	19.66	37.71	40.79	41.28	34.82	14.88	17.29	25.76	7.33
RAMMA1000510	8.43	9.68	12.58	8.8	8.42	5.47	7.38	4.77	5.1
RAMMA1000515	16.44	20.18	30.29	25.14	20.54	20.81	15.29	9.87	15.02
RAMMA1000516	5.05	38.96	14.92	3.28	5.69	7.88	10.33	1.91	1.98
RAMMA1000522	27.99	43.17	68.38	38.83	18.51	22.66	24.89	20.61	23
RAMMA1000524	5.97	11.73	10.41	4.38	10.1	5.92	6.73	2.46	2.78
RAMMA1000528	3.99	4.53	7.16	4.83	4.31	6.82	4.96	1.75	1.88
RAMMA1000534	20.75	20.93	19.83	14.18	15.4	10.89	14.95	12.85	11.39
RAMMA1000541	7.6	8.83	11.36	1.44	4.17	2.11	2.2	1.84	1.88
RAMMA1000550	2.63	1.99	8.54	2.03	3.94	1.32	3.32	0.71	2.83
RAMMA1000556	761.19	1381.9	1153	543.64	1435.2	1032.9	1229.3	435.34	855.28
RAMMA1000559	8.09	5.3	11.69	2.87	3.18	2.37	2.27	0.81	3.09
RAMMA1000565	3.15	5.75	13.35	1.46	4.28	0.82	2.18	0.1	1.13

40 / 157

【0864】

【表393】

2519

2520

RAM1000587	2.83	9.61	12.53	4.59	4.96	3.08	3.91	1.78	2.49
RAM1000588	14.4	23.94	41.43	6.77	7.53	16.68	14.73	20.93	24.41
RAM1000589	342.82	404.05	589.09	206.24	397.1	307.2	196.5	234.99	295.12
RAM1000590	3.44	4.94	9.73	2.95	5.37	2.97	1.64	1.64	2.31
RAM1000591	10.38	9.76	14.3	7.6	8.3	4.54	9.1	4.55	7.57
RAM1000592	9.1	16.85	23.22	12.88	13.24	10.32	9.64	5.62	11
RAM1000593	24.67	34.19	52.67	17.16	22.85	23.65	15.05	11.26	16.03
RAM1000594	25.66	32.59	51.71	40.6	49.12	36.4	18.51	9.64	20.88
RAM1000595	10.5	17.98	38.06	21.57	17.46	15.03	12.1	5.53	10.65
RAM1000596	8.03	8.9	14.02	3.22	4.88	8.25	3.07	3.05	5.3
RAM1000597	3.76	3.87	7.3	1.87	2.17	1.53	2.62	0.72	2.53
RAM1000598	8.91	9.35	17.48	7.74	5.84	6.31	7.11	2.55	5.81
RAM1000599	10.09	7	22.58	5.69	12.68	6.11	4.8	4.88	3.1
RAM1000600	18.37	28.27	42.33	24.96	17.41	18.13	16.86	13.76	19.28
RAM1000601	5.51	4.72	11.4	2.5	4.83	2.59	2.52	1.82	1.65
RAM1000602	3.19	6.18	12.95	3.95	4.29	3.37	3.1	1.01	2.69
RAM1000603	3.08	4.28	9.78	5.68	3.5	2.91	2.33	0.32	2.12
RAM1000604	3.16	4.8	8.96	4.43	4.19	14.5	3.27	1.73	4.49
RAM1000605	53.05	63.8	128.68	68.08	63.84	76.25	48.7	34.2	32.4
RAM1000606	7.49	6.59	12.24	7.94	7.8	10.34	5.92	2.82	1.26
RAM1000607	4.63	4.12	7.44	5.71	2.66	4.08	1.99	2.69	1.6
RAM1000608	18.71	22.51	38.34	17.17	14.96	18	10.64	12.09	11.81
RAM1000609	8.69	17.36	17.2	8.58	6.25	8.92	3.77	5.35	2.7
RAM1000610	23.14	46.99	61.08	27.38	33.9	32.87	41.85	23.92	33.78
RAM1000611	3.03	3.5	9.1	2.2	6.51	1.81	2.38	2.13	1.46
RAM1000612	4.89	6.68	9.53	5.24	3.91	6.43	9.4	3.5	1.93
RAM1000613	127	118.39	232.08	362.26	175.38	245.8	112.44	89.34	106.16
RAM1000614	42.17	56.4	86.57	28.1	10.46	18.45	7.15	8.3	5.57
RAM1000615	6.35	5.08	11.75	8.34	4.22	4.13	4.18	2.86	3.12
RAM1000616	7.63	9.04	16.29	12.54	11.44	15.76	4.16	3.21	6.89
RAM1000617	2.37	2.75	7.2	4.27	3.94	4.85	2.38	1.14	1.76
RAM1000618	7.11	13.17	13.45	8.85	13.62	11.39	5.11	2.82	4.49
RAM1000619	5.79	9.88	8.48	5.04	7.37	7.72	4.37	2.73	5.55
RAM1000620	13.18	25.83	23.83	10.07	8.95	6.62	3.02	4.1	4.74
RAM1000621	21.05	28.79	38.76	24.92	18.84	21.25	20.38	21.53	20.87
RAM1000622	7.11	5.26	13.51	4.07	3.72	3	2.71	1.31	2.37
RAM1000623	56.24	56.67	100.19	108.88	65.38	82.83	53.41	37.73	50.75
RAM1000624	83.95	62.78	122.37	158.83	112.68	140.59	85.25	48.89	63.88
RAM1000625	13.21	20.89	24.92	19.74	19.6	18.49	13.07	8.64	14.74
RAM1000626	4.13	6	9.07	5.99	9.97	7.98	4.18	1.33	3.64
RAM1000627	12.22	18.56	20.77	15.57	15.03	11.37	17.6	9.79	15.79
RAM1000628	4.23	6.43	10.91	7.98	3.73	6.91	3.13	3.72	3.43
RAM1000629	4.58	4.07	5.19	5.1	2.18	3.84	1.48	1.8	2.72
RAM1000630	27.47	31.39	43.2	35.52	27.09	30.72	11.08	15.65	19.91
RAM1000631	11.43	16.18	20.32	9.83	9.84	10.4	18.64	15.89	17.27
RAM1000632	5.14	6.4	11.37	3.9	6.11	4.85	2.11	2.43	3.5
RAM1000633	2.95	3.93	9.11	2.68	4.26	3.62	2.53	1.16	2.48
RAM1000634	3.63	4.94	7.73	2.22	4.21	3.15	2.53	1.58	2.63
RAM1000635	9.18	9.76	16.46	16.7	13	12.37	10.17	5.29	10.03
RAM1000636	17.77	21.89	38.71	25.58	19.94	21.44	12.48	15.54	20.02
RAM1000637	6.79	10.56	9.64	3.73	4.03	3.96	1.97	4.15	4.68
RAM1000638	4.08	4.2	7.68	1.53	2.88	3.57	1.98	1.33	1.45
RAM1000639	4.12	5.84	14.4	7.07	2.94	5.07	2.8	2.71	4.1
RAM1000640	26.15	27.77	36.92	48.47	26.15	33.74	22.66	12.7	18.46
RAM1000641	38.93	73.17	60.3	15.58	59.91	40.66	16.6	19.11	12.75
RAM1000642	10.71	16.72	21.46	6.1	15.65	13.14	10.25	9.83	6.19
RAM1000643	5.28	6.96	8.31	4.59	9.33	5.12	3.95	1.7	2.78
RAM1000644	12.93	21.78	27.25	9.87	8.6	8.24	5.41	7.95	18.83
RAM1000645	14.83	19.2	21.42	14.77	17.9	18.09	9.03	8.51	18.28
RAM1000646	10.63	12.91	13.99	8.66	4.6	7.82	5.32	3.01	7.59
RAM1000647	10.59	10.38	19.17	11.33	9.88	8.79	5.57	3.43	6.21
RAM1000648	8.09	8.16	13.72	6.5	8.85	7.38	2.82	2.22	3.59
RAM1000649	22.11	37.42	97.78	18.99	20.99	20.43	27.71	18.27	23.96
RAM1000650	1.64	5.26	13.69	2.27	7.88	1.6	3.8	1.08	1.8

41 / 157

【0865】

【表394】

2521

2522

RAM1000824	52.76	82.46	54.88	22.76	88.37	48.11	53.27	21.38	20.22				
RAM1000827	3.29	5.2	12.5	6.91	5.04	2.83	2.54	1.96	3.02				
RAM1000831	31.08	36.42	39.98	20.36	22.38	25.87	17.62	21.92	26.79	*	*	-	-
RAM1000836	4.58	6.92	8.85	2.9	2.71	3.02	3.13	2.4	1.8	*	*	-	-
RAM1000839	4.7	5.1	9.24	1.82	2.67	4.2	2.02	1.34	1.48	*	*	-	-
RAM1000841	4.43	2.78	9.88	2.71	2.83	2.07	3.01	0.81	2.24				
RAM1000842	6.22	8.19	14.72	3.25	5.48	4.22	5.77	2.92	7.82				
RAM1000843	10.02	23.1	28.68	14.54	17.26	10.11	4.69	3.85	6.21	*		-	-
RAM1000845	2.67	2.89	10.34	0.88	3.81	2.04	1.66	1.53	3.4				
RAM1000851	40.29	50.53	100.88	64.55	45.92	39.69	42.2	16.12	64.01				
RAM1000854	3.86	5.86	8.12	2.91	1.76	3.58	1.11	0.53	1.24	*		-	-
RAM1000855	91.57	117.96	169.35	219.71	138.54	171.88	95.86	65.57	93.06				
RAM1000855	3.13	2.69	5.73	1.9	1.05	2.68	1.82	0.46	1.81				
RAM1000859	3.59	5.65	10.37	1.78	1.83	1.2	3.39	1.89	1.24				
RAM1000882	8.58	10.61	23.89	5.34	11.70	7.43	5.16	1.27	2.11				
RAM1000883	17.14	20.47	45.55	14.38	24.31	16.09	24.81	12.68	16.34				
RAM1000885	3.2	5.96	11.12	4.18	4.78	11.8	2.48	0	2.86				
RAM1000887	12.55	18	37.87	18.03	16.36	15.87	9.25	8.53	14.8				
RAM1000875	197.47	247.19	107.15	151.05	163.7	159.54	127.81	148.8	137.51	*	*	-	-
RAM1000876	4.29	6.54	6.75	2.84	2.75	3.43	2.51	1.58	2.61	*	*	-	-
RAM1000877	2.84	3.61	4.96	2.05	1.13	1.68	2.93	0.77	2.2	*		-	-
RAM1000878	5.22	9.31	15.62	6.72	6.38	5.06	10.1	4.12	5.11				
RAM1000880	13.16	16.7	32.88	12.81	12.85	8.92	10.54	7.92	11.52				
RAM1000881	6.97	11.85	26.23	10.06	11.84	7.67	8.29	6.16	7.32				
RAM1000883	11.76	21.66	31.01	10.87	18.82	18.1	18.77	10.78	18.11				
RAM1000887	16.89	26.07	44.99	26.44	17.63	21.31	17.58	14.36	19.92				
RAM1000888	4.54	4.74	8.28	5.15	4.11	4	3.9	1.83	1.2				
RAM1000905	19.81	46.58	24.87	13.58	12.04	12.03	17.95	11.63	11.81				
RAM1000906	13.7	9.74	18.2	5.83	6.94	4.39	4.39	2.34	1.82	*	*	-	-
RAM1000908	8.3	13.83	18.11	10.1	14.32	5.63	5.08	3.21	3.69	*		-	-
RAM1000911	13.12	16.04	26.46	13.81	12.11	7.57	12.29	6.96	12.25				
RAM1000914	2.41	3.47	11.09	1.11	2.25	1.29	1.59	0	0.87				
RAM1000920	68.92	152.1	139.08	77.3	186.8	94.88	122.37	27.22	25.46				
RAM1000921	3.68	5.93	12.87	7.11	6.12	6.97	3.45	2.44	2.82				
RAM1000931	35.48	29.12	32.19	9.74	14.81	15.82	7.85	20.92	4.78	**	*	-	-
RAM1000940	16.07	19.99	34.79	11.62	14.07	12.51	10.88	8.84	14.41				
RAM1000941	38.94	28.8	55.32	14.85	48.89	25.1	44.15	31.17	20.09				
RAM1000942	19.99	16.82	58.05	28.3	25.6	20.38	16.91	16.31	24.51				
RAM1000943	14.05	10.94	26.95	6.92	4.75	8.24	3.89	2.4	1.4	*		-	-
RAM1000952	3.98	5.7	16.6	3.37	4.87	3.51	7.54	1.89	3				
RAM1000956	5.62	5.18	14.83	5.99	5.03	6.18	4.85	2.51	5.15				
RAM1000957	0.61	0.47	2.13	0.38	0.27	1.52	0.16	0	0				
RAM1000962	26.82	26.6	21.85	10.69	8.37	7.64	7.43	8.92	3.07	**	**	-	-
RAM1000966	13.96	15.32	16.62	3.18	4.77	5.42	5.48	5.55	2.07	**	**	-	-
RAM1000968	3.62	3.17	8.04	2.18	4.04	1.69	2.56	0.98	1.54				
RAM1000972	109.05	38.4	188	38.59	38.3	50.84	51.28	65.81	23.12				
RAM1000973	4.24	3.8	8.43	1.16	1.43	2.19	3.18	1.23	1.88				
RAM1000975	4.83	5.04	15.94	8.44	6.12	4.99	5.27	1.93	5.12				
RAM1000976	10.4	12.69	31.13	7.68	10.7	11.37	8.68	4.42	4.88				
RAM1000979	15.33	18.37	49.49	29.29	22.41	26.44	15.03	8.9	16				
RAM1000986	11.83	20.17	19.1	14.27	9.07	14.47	9.64	5.8	3.91	*		-	-
RAM1000987	35.85	39.55	84.08	87.43	82.8	61.8	46.73	26.89	38.89				
RAM1000988	12.53	13.76	42.26	22.92	17.83	17.38	12.36	7.34	12.64				
RAM1000994	6.93	10.09	26.01	16.14	13.64	12.99	6.56	3.79	6.83				
RAM1000998	3.85	8.09	10.95	3.78	4.4	5.85	1.97	0.73	2.47	*		-	-
RAM1001003	11.87	14.9	34.42	24.16	21.08	20.65	13.23	10.02	16.87				
RAM1001007	4	4.13	8.2	3.14	8.73	3.28	4.14	1.49	4.79				
RAM1001008	20.12	23.78	44.26	26.75	20.79	27.13	18.57	19.08	24.22				
RAM1001013	9.3	12.99	14.08	13.97	3.4	7.56	6.85	5.48	5.5	*		-	-
RAM1001014	3.74	2.89	7.41	3.88	5.49	8.98	4.67	4.19	1.31				
RAM1001021	58.82	76.19	93.74	67.88	54.85	51.75	72.35	45.95	58.5				
RAM1001024	88.7	92.15	131.2	139.18	94.51	99.4	87.88	68.05	108.24				
RAM1001025	69.9	102.4	41.86	70.13	59.68	30.75	22.6	25.23	39.06				
RAM1001028	80.18	111.58	110	73.98	84.44	72.45	97.15	91.89	100.75				

42 / 157

【0866】

【表395】

2523

2524

HAIRMA1001030	7	7	11.72	12.89	11.66	7.79	10.11	7.26	6.15	7.3
HAIRMA1001035	39.22	53.03	74.6	50.1	27.34	38.78	23.51	81.74	38.44	
HAIRMA1001036	3.31	5.83	5.4	5.82	5.35	4.3	4.37	2.71	3.74	
HAIRMA1001037	24.23	31.03	53.22	27.23	25.26	31.34	28.91	17.23	26.48	
HAIRMA1001038	12.28	20.58	21.05	12.13	14.71	11.55	14.52	10.1	18.77	
HAIRMA1001041	1.64	2.15	6.75	4.24	4.69	7.8	3.02	1.22	2.08	
HAIRMA1001043	1.39	2.46	4.67	3.06	4.27	3.32	4.35	1.52	2.3	
HAIRMA1001050	1.33	4.39	4.83	3.8	5.59	6.14	4.27	1.05	2.57	
HAIRMA1001054	15.75	24.03	50.38	32.02	21.79	27.11	13.83	16.21	20.46	
HAIRMA1001058	2.19	4.55	5.53	3.65	3.25	5.05	2.54	10.99	3.33	
HAIRMA1001066	5.02	4.16	8.84	5.86	2.85	3.72	1.82	2.25	0.88	
HAIRMA1001067	17.58	21.79	31.08	14.83	16.92	18.84	15.29	17.34	18.26	
HAIRMA1001072	2.53	1.68	3.3	1.93	1.68	2.41	2.3	1.26	2	
HAIRMA1001073	0.84	2.27	6.85	1.01	1.41	5.24	0.75	4.07	1.61	
HAIRMA1001074	24.28	39.17	47.61	31.86	28.12	24.54	31.91	23.64	32.81	
HAIRMA1001075	1.48	5.48	8.44	3.98	4.91	2.99	2.44	2.09	3.72	
HAIRMA1001076	21.72	28.12	44.96	39.39	16.5	21.38	18.07	17.35	25.34	
HAIRMA1001080	12.04	17.72	15.6	14.28	12.08	16.7	6.39	6.97	11.08	
HAIRMA1001082	4.69	9.29	12.12	6.84	2.79	4.23	4.1	1.12	3.78	
HAIRMA1001091	18.73	23.95	30.14	19.81	16.48	17.71	21.26	12.1	23.8	
HAIRMA1001092	7.71	11.13	20.6	7.44	8.17	8.28	7.18	5.4	7.58	
HAIRMA1001094	11.74	18.72	15.33	14.77	10.66	23.71	14.53	12.3	17.19	
HAIRMA1001095	45.26	57.55	68.73	41.25	47.01	32.75	30.09	24.22	28.58	
HAIRMA1001110	7.59	14.4	12.56	8.84	9.45	3.97	9.82	4.24	0.9	
HAIRMA1001126	37.56	44.85	64.49	49.13	24.88	37.49	24.53	36.31	49.73	
HAIRMA1001133	6.76	9.31	8.87	3.28	3.01	4.08	2.84	5.37	6.55	
HAIRMA1001139	2.77	14.13	10.95	4.54	4.49	4.1	6.16	2.44	5.84	
HAIRMA1001141	6.69	8.3	15.1	2.8	4.24	4.06	2.47	2.3	5.32	
HAIRMA1001142	3.83	3.21	9.91	1.84	2.97	1.81	3.49	1.28	2.11	
HAIRMA1001145	2.22	12.54	14.95	2.6	3.58	6.73	2.4	2.78	3.58	
HAIRMA1001150	10.97	15.54	29.9	12.2	12.68	9.93	10.31	8.5	12.03	
HAIRMA1001154	5.02	9.05	11.37	4.94	8.38	40.3	3.8	2.43	3.42	
HAIRMA1001159	6.91	7.68	15.13	5.63	3.35	4.72	2.34	2.78	4.72	
HAIRMA1001161	109.98	158.02	158.22	189.81	141.37	144.88	81.63	94.69	118.15	
HAIRMA1001162	7.19	6.58	7.65	7.64	6.6	6.96	4.02	2.28	5.78	
HAIRMA1001163	2.37	3.1	6.4	4.16	1.93	2.21	2.1	2.11	4.1	
HAIRMA1001166	3.02	2.01	9.14	2.7	2.02	2.05	2.94	0.77	3.58	
HAIRMA1001189	17.92	24.89	44.3	19.97	22.88	16.28	24.9	14.32	18.92	
HAIRMA1001191	10.11	15.9	19.2	7.11	13.08	8.79	15.3	10.81	18.52	
HAIRMA1001198	4.68	8.28	11.86	3.48	7.28	5.15	6.52	2.84	2.13	
HAIRMA1001202	30.38	37.64	84.46	54.02	29.63	36.02	18.16	17.9	40.49	
HAIRMA1001203	8.99	12.77	20.34	10.56	11.45	9.86	10.07	8.3	11.49	
HAIRMA1001205	14.27	17.3	31.28	14.44	9.47	13.95	15.09	7.44	16.20	
HAIRMA1001206	3.37	1.85	7.28	0.99	1.62	0.23	1.58	0.63	2.48	
HAIRMA1001215	12.57	20.58	26.01	4.38	17.56	6.44	10.82	1.62	8.27	
HAIRMA1001220	12.12	14.97	20.97	18.99	11.55	15.4	10.87	5.54	12.63	
HAIRMA1001222	13.14	15.27	54.43	12	15.72	10.61	17.81	13.51	16.5	
HAIRMA1001223	5.47	11.04	23.87	8.51	6.78	8.32	9.09	3.98	9.79	
HAIRMA1001232	5.9	6.27	10.44	10.05	4.68	7.4	3.45	2.12	5.69	
HAIRMA1001234	78.47	62.1	108.68	184.09	22.14	88.22	44.43	81.83	25.73	
HAIRMA1001237	4.23	3.61	5.41	2.39	3.8	1.85	3.83	2.61	6.33	
HAIRMA1001243	28.51	33.45	37.19	22.72	19.97	20.13	13.4	8.73	8.38	
HAIRMA1001244	2.39	3.23	6.47	2.05	2.61	0.89	2.52	0.99	1.85	
HAIRMA1001249	32.97	38.57	66.31	31.72	48.81	37.57	44.31	25.2	27.49	
HAIRMA1001256	11.6	15.37	22.1	10.86	18.44	20.14	15.27	12.69	11.49	
HAIRMA1001259	3.58	4.73	9.83	2.71	2.58	1.74	5.81	0.4	3.31	
HAIRMA1001260	15.61	19.65	19.14	11.02	7.29	22.35	9.03	6.24	10.39	
HAIRMA1001282	8.3	48.53	16.27	15.78	7.11	6.47	9.48	5.41	7.71	
HAIRMA1001288	7.28	6.86	12.6	6.74	6.05	4.12	3.69	3.37	4.04	
HAIRMA1001291	4.99	4.98	6.83	2.12	3.55	3.26	4.68	2.4	2.94	
HAIRMA1001274	197.38	283.08	348.26	54.58	91.98	70.74	33.25	18.31	25.39	
HAIRMA1001280	10.76	14.33	28.16	10.36	11.35	10.7	14.52	8.71	14.89	
HAIRMA1001283	6.41	3.31	10.89	0.34	3.46	2.07	2.94	1.3	1.97	
HAIRMA1001284	6.24	6.15	16.76	7	6.77	4.3	3.05	1.61	2.99	

43 / 157

2525

2526

RAM1A1001280	29.08	30.31	36.62	8.75	7.01	14.48	10.18	18.99	22.31	**	-
RAM1A1001289	8.87	7.86	10.4	8.73	7.53	9	7.31	3.62	7.73		
RAM1A1001292	24.97	44.07	66.64	9.34	36.59	35.11	13.77	28.43	10.4		
RAM1A1001295	4	4.17	8.01	2.8	3.49	2.38	2.75	2.82	4.19		
RAM1A1001298	7.53	9.89	10.46	2.88	6.46	2.62	5.34	3.84	1.86	*	-
RAM1A1001305	3.74	4.35	13.81	2.61	4.06	3.12	3.76	6.53	2.5		
RAM1A1001308	10.42	15.14	31.52	12.87	13.99	9.19	10.23	3.88	9.86		
RAM1A1001310	2.92	5.46	13.25	3.59	3.89	1.96	3.03	0.43	1.21		
RAM1A1001322	2.17	2.64	6.93	3.3	2.91	2.5	1.12	1.37	2.03		
RAM1A1001324	4.53	3.69	6.65	3.83	2.78	3	2.85	1.6	1.68	*	-
RAM1A1001330	4.9	5.1	12.68	7.84	4.19	4.82	3.35	2.64	3.18		
RAM1A1001333	4.95	3.16	7.58	2.58	3.62	3.33	2.38	2.25	1.89		
RAM1A1001334	31.92	25.86	49.45	18.83	15.13	18.08	17.92	22.66	13.99	*	-
RAM1A1001337	4.26	3.84	13.74	3.62	4.64	5.34	4	3.29	3.18		
RAM1A1001341	4.05	4.98	11.41	4.38	5.21	4.83	5.24	4.52	3.64		
RAM1A1001343	2.73	5.65	10.8	3.22	4.68	4.2	3.94	1.64	3.95		
RAM1A1001344	2.25	2.47	7.73	4.27	4.38	3.62	2.73	1.3	3.12		
RAM1A1001345	5.84	8.71	17.37	10.65	9.35	8.33	6.74	3.26	6.06		
RAM1A1001383	8.99	8.47	18.39	14.27	8.27	9.89	7.45	4.12	8.12		
RAM1A1001385	25.28	28.84	63.23	32.5	23.88	25.62	24.7	25.15	17.89		
RAM1A1001386	15.94	20.37	30.49	14.54	24.62	28.46	14.45	46.08	11.25		
RAM1A1001387	14.75	18.37	41.81	21.11	18.82	18.79	17.96	11.5	21.37		
RAM1A1001401	2.94	3.12	6.66	1.21	2.62	3.02	2.45	0.78	2.89		
RAM1A1001408	1.5	3.57	7.4	2.81	1.88	3.96	5.62	0.78	2.48		
RAM1A1001411	6.08	3.93	9.81	9.48	8.09	8.29	3.35	4.18	4.81		
RAM1A1001414	6.51	7.32	14.7	23.79	6.88	10.87	5.51	4.82	6.45		
RAM1A1001415	3.95	8.97	6.68	5.52	5.88	8.18	6.05	2.79	4.69		
RAM1A1001418	8.77	9.1	12.47	12.83	10.08	11.32	13.03	7.4	7.63		
RAM1A1001419	7.73	7.93	11.48	9.88	8.73	10.35	9.18	5.19	8.18		
RAM1A1001420	5.14	4.76	6.87	10.23	8.22	6.68	5.34	3.95	4.54		
RAM1A1001426	5.19	7.71	9.63	5.55	8.11	6.73	7.6	4.91	8.87		
RAM1A1001428	3.64	3.18	7.82	8.62	3.95	5.09	3.35	1.49	3.18		
RAM1A1001432	12.35	13.68	21.83	18.83	11.53	18.33	8.1	11.28	14.53		
RAM1A1001435	29.05	30.02	44.24	21.04	23.86	21.39	14.97	14.62	18.51		
RAM1A1001442	8.94	7.75	11.08	5.63	5.57	7.7	8.58	4.54	7.44		
RAM1A1001446	28.48	31.11	44.32	20.83	23.27	24.39	8.85	5.14	8.85	**	-
RAM1A1001450	11.55	16.7	15.65	12.27	11.07	10.5	14.48	8.46	14.95		
RAM1A1001452	4.88	7.91	8.35	9.2	4.08	8.24	7.72	3.93	8.05		
RAM1A1001485	8.92	12.52	15.39	11.83	13.04	10.87	15.92	7.42	13.08		
RAM1A1001476	132.51	144.85	285.89	418.87	194.88	350.68	82.5	94.08	100.24		
RAM1A1001478	18.64	21.54	45.82	21.72	20.69	19.34	14.71	16.9	28		
RAM1A1001479	11.41	20.27	19.2	13.74	10.72	9.98	13.4	11.81	15.33		
RAM1A1001487	48.7	29.91	87.89	41.22	44.32	54.61	29.8	35.15	13.98		
RAM1A1001498	4.35	2.98	8.82	3.07	3.33	1.89	1.92	4.02	3.09		
RAM1A1001501	4	5.27	10.82	3.92	4.81	6.08	4.06	2.78	3.9		
RAM1A1001502	3.5	3.68	8.07	3.61	4.35	4.14	3.43	1.95	7.01		
RAM1A1001510	11.45	20.55	16.62	13.12	12.05	13.04	13.39	7.86	10.37		
RAM1A1001522	4.88	10.87	13.99	8.11	4.33	5	2.48	3.78	5.04		
RAM1A1001529	59.97	69.9	94.67	30.63	48.88	57.69	31.84	37.9	51.83	*	-
RAM1A1001532	28.38	37.51	51.48	23.78	28.62	23.49	28.8	24.11	41.98		
RAM1A1001533	19.86	8.23	28.5	6.01	12.09	8.61	7.93	4.83	6.28		
RAM1A1001534	11.45	13.72	20.65	6.58	8.79	8.81	8.6	5.71	8.05		
RAM1A1001535	10.07	15.11	17.7	7.42	8.38	7	2.75	2.21	3.48	*	-
RAM1A1001547	5.79	10.79	16.77	6.18	7.84	8.73	7.59	4.04	7.64		
RAM1A1001551	1.79	3.35	10.09	2.36	3.76	2.54	3.47	0.46	2.39		
RAM1A1001562	5.44	7.64	12.53	5.97	3.18	5.89	2.71	3.74	6.78		
RAM1A1001575	19.87	31.06	32.95	20.06	20.37	18.83	18.83	18.85	22.11		
RAM1A1001576	4.97	13.68	9.34	4.65	6.28	3.49	1.76	5.49	4.18		
RAM1A1001584	17.12	20.22	32.05	43.71	22.66	30.09	21.3	15.83	24.89		
RAM1A1001588	3.44	2.63	9.42	2.75	2.62	1.44	2.63	0.98	2.02		
RAM1A1001590	4.74	3.32	9.43	0.85	2.72	2.98	5.93	1.26	2.67		
RAM1A1001599	6.91	10.35	19.33	5.33	6.4	5.4	7.64	5.96	7.44		
RAM1A1001800	12.68	24.27	25.25	19.57	15.9	18.81	21.18	12.02	17.72		
RAM1A1001804	20.93	20.87	41.04	17.28	15.67	6.48	4.43	10.16	15.17		

44 / 157

2527

2528

RAMMA1001605	37.25	58.33	44.14	13.41	28.09	29.22	7.5	16	15.18	*	**	-	-
RAMMA1001609	2.18	4.53	5.97	1.87	1.39	1.8	4.17	2.27	1.51				
RAMMA1001614	10.91	17.85	21.09	12.42	12.51	13.02	18.85	9.92	19.29				
RAMMA1001615	1.95	3.17	11.55	2.77	2.24	1.35	2.33	1.48	2.05				
RAMMA1001619	11.51	18.15	15.55	5.34	13.88	8.59	8.19	7.08	5.14	*		-	
RAMMA1001620	4.59	6.42	11.48	1.77	3.88	3.43	3.11	1.71	4				
RAMMA1001623	48.38	31	53.81	63.16	42.72	47.15	31.15	25.06	25.68				
RAMMA1001626	23.59	24.5	38.75	33.26	30.12	18.9	7.93	11.83	22.41				
RAMMA1001627	23.89	43.37	29.47	17.73	21.82	12.9	6.43	10	7.98	*		-	
RAMMA1001630	9.27	13.78	11.56	7.25	5.83	6.82	2.81	4.06	2.66	*	**	-	
RAMMA1001633	3.1	2.16	7.03	2.23	2.28	2.15	0.8	0.79	2.49				
RAMMA1001634	1.53	6.28	9.01	3.76	4.19	2.88	3.9	2.94	2.79				
RAMMA1001635	1.28	4.64	15.97	2.28	2.56	1.53	6.22	1.65	2.6				
RAMMA1001649	1.05	5.72	15.55	3.26	4.9	2.56	4.28	1.39	3.84				
RAMMA1001654	4.23	5.9	13.55	3.28	5.84	3.8	3.83	1.4	2.71				
RAMMA1001660	39.09	42.24	88.43	56.58	52.47	44.31	41.47	39	60.86				
RAMMA1001683	8.52	11.06	11.71	10.2	6.57	5.19	7.98	6.64	8.33				
RAMMA1001670	5.91	7.75	6.72	4.07	2.57	5.22	5.32	3.26	4.27	*	*	-	
RAMMA1001671	58.81	62.54	66.07	56.8	37.97	53.72	50.14	43.86	58.86				
RAMMA1001679	8.69	13.86	10.25	4.05	5.47	3.58	4.48	7.43	6.03				
RAMMA1001683	15.8	21.19	31.72	12.85	17.62	10.3	18.14	18.49	14.57				
RAMMA1001686	4.11	3.45	11.77	4.88	4.36	4.4	4.89	1.58	2.69				
RAMMA1001688	4.98	5.41	15.23	3.79	4.7	4.42	2.85	1.45	3.58				
RAMMA1001689	28.44	25.85	32.18	7.85	17.39	15.22	8.44	8.89	20.79	*	*	-	
RAMMA1001692	6.97	3.79	10.24	3.42	9.87	3.1	3.28	4.26	5.93				
RAMMA1001711	4.75	4.68	8.45	3.21	3.28	2.22	2.67	8.04	0.88				
RAMMA1001715	4.22	4.73	9.43	3.9	2.04	3.47	3.76	4.11	2.61				
RAMMA1001730	3.39	2.22	8.02	1.72	1.43	1.79	1.71	1.46	1.88				
RAMMA1001735	8.81	9.98	22.83	12.18	11.12	10.37	6.84	5.43	7.1				
RAMMA1001740	2.28	4.25	11.58	2.64	4.24	1.86	2.43	1.41	3.14				
RAMMA1001743	10.52	18.5	19.92	5.55	7.23	6.37	8.45	1.87	3.83	*	*	-	
RAMMA1001744	4.59	3.07	5.75	3.88	4	4.18	2.05	2.19	3.63				
RAMMA1001745	12.82	11.29	21.77	12.13	11.56	10.5	13.92	9.55	13.74				
RAMMA1001751	11.11	8.28	15.68	3.47	4.56	3.38	2.93	2.2	1.66	*	*	-	
RAMMA1001752	5.63	3.02	9.77	2.83	5.25	2.56	2.97	2.42	0.87				
RAMMA1001754	23.39	32.69	51.88	31.96	35.9	28.06	34.24	24.41	25.71				
RAMMA1001757	5.44	8.41	15.4	4.99	7.7	6.28	7.64	3.88	6.12				
RAMMA1001760	2.58	5.03	10.84	4.07	3.09	11.35	3.87	1.47	1.67				
RAMMA1001764	1.11	5.18	12.27	3.88	5.02	2.65	4.09	8.44	2.15				
RAMMA1001767	3.1	2.25	5.04	3.83	2.38	7.31	2.64	1.02	1.81				
RAMMA1001768	11.92	10.35	26.44	10.8	12.51	10.66	12.91	3.89	9.89				
RAMMA1001769	14.67	15.11	32.18	14.99	14.79	13.17	17.74	10.68	13.05				
RAMMA1001771	5.81	5.01	8.7	5.52	6.04	4.42	3.1	4.15	2.2				
RAMMA1001773	40.73	48.88	88.4	41.31	38.76	36.3	25.57	21.1	22.63				
RAMMA1001778	15.44	10.48	54.31	14.07	14.31	18.11	18.19	12.85	7.66				
RAMMA1001783	12.5	12.45	19.56	11.82	10.75	17.89	7.63	5.83	3.14				
RAMMA1001785	4.08	3.99	8.35	2.35	3.38	3.07	3.75	1.41	2.68				
RAMMA1001788	6.1	6.55	20.74	18.01	10.68	15.67	7.37	4.2	7.06				
RAMMA1001790	1.31	0.72	1.87	1.36	1.35	1.32	1.67	0.59	1.38				
RAMMA1001800	2.19	1.85	2.86	3.44	3.43	3.44	1.48	0.44	1.31	**	*	+	-
RAMMA1001804	9.74	13.74	28.09	23.09	17.83	16.06	15.83	9.04	8.32				
RAMMA1001806	15.42	19.2	24.15	21.08	18.33	17.7	13.08	11.51	12.4				
RAMMA1001812	2.68	2.87	5.53	5.25	2.66	2.24	1.76	1.16	2				
RAMMA1001815	0.82	1.33	4.14	2.35	1.64	0.9	2.33	0	1.04				
RAMMA1001817	7.16	8.39	26.7	8.94	11.15	9.35	10.8	5.97	8.95				
RAMMA1001818	11.74	18.24	23.68	11.92	10.04	13.23	8.57	10.05	9.02				
RAMMA1001819	15.72	24.34	49.48	17.35	11.3	16.18	12.64	12.63	16.41				
RAMMA1001820	22.62	19.64	37.64	42.37	24.18	31.88	15.13	16.66	19.01				
RAMMA1001824	8.99	12.77	18.28	12.69	14.09	15.06	12.72	8.76	11.85				
RAMMA1001832	7.22	10.73	12.88	9.65	7.55	9.34	6.24	4.93	7.88				
RAMMA1001836	9.42	23.13	19.47	21.69	15.29	10.85	11.8	5.71	11.11				
RAMMA1001837	3.6	4.29	8.23	5.89	6.7	8.38	4.82	1.11	4.3				
RAMMA1001848	16.38	27.48	23.83	25.7	38.67	49.58	24.69	13.45	27.7				
RAMMA1001850	4.58	4.63	5.84	5.14	2.91	3.96	4.34	2.05	2				

45 / 157

【0869】

【表398】

2529

2530

SHAMMA1001851	3.04	3.08	4.42	3.11	3.54	2.54	2.52	0.56	2.38
SHAMMA1001852	4.14	3.26	11.19	3.61	3.33	5.4	1.88	2.12	2.02
SHAMMA1001854	1.51	2.81	8.9	2.97	1.88	2.18	2.29	1.4	2.53
SHAMMA1001858	19.51	21.77	35.8	19.05	18.81	18.88	15.59	11.41	20.08
SHAMMA1001854	13.26	17.9	34.07	19.43	19.24	18.98	19.91	10.97	23.39
SHAMMA1001868	36.41	51.95	37.39	27.02	32.08	24.02	38.47	28.02	40.25
SHAMMA1001874	10.43	12.88	12.56	15.09	16.33	15.08	13.47	8	7.94
SHAMMA1001878	8.74	13.09	17.2	10.72	7.64	9.1	8.72	5.47	7.12
SHAMMA1001880	4.07	4.28	5.15	5.16	2.51	4.87	2.78	4.1	3.13
SHAMMA1001885	11.53	15.59	29.89	11.02	9.99	7.12	10.84	10.3	12.47
SHAMMA1001890	18	26.55	27.35	18.41	24.49	18.88	18.85	18.1	22.42
SHAMMA1001892	9.74	13.29	21.33	8.08	11.26	12.34	13.42	8.15	13.42
SHAMMA1001901	5.91	10.54	9.81	8.07	8.49	8.27	5.4	3.05	2.51
SHAMMA1001907	2.81	3.52	7.04	35.82	7.01	7.16	2.77	1.6	2.87
SHAMMA1001908	8.52	7.37	12.33	5.04	8.79	5.83	5.18	1.89	2.73
SHAMMA1001919	28.87	31.67	48.08	30.53	41.28	68.58	17.81	21.05	28
SHAMMA1001931	5.05	3.85	9.8	4.52	3.24	2.79	3.04	5.51	2.52
SHAMMA1001937	14.28	19.97	17.05	11.88	9.87	18.45	11.3	11.49	14.97
SHAMMA1001951	10.03	11.71	22.7	11.88	8.84	10.7	11.47	12.01	12.62
SHAMMA1001958	993.12	926.97	1643.3	1427.1	1487.9	1519.7	1082.9	902.24	975.15
SHAMMA1001957	5.28	8.18	19.85	8.48	7.2	6.6	6.19	4.02	7.97
SHAMMA1001960	4.45	5.22	11.99	3.94	5.66	4.15	4.01	2.89	2.74
SHAMMA1001963	9.87	14.08	20.88	14.68	14.73	14.4	8.07	8	3.43
SHAMMA1001969	11.23	14.81	19.75	14.09	10.96	16.21	10.31	8.8	19.07
SHAMMA1001970	4.02	5.46	5.52	3.48	4.15	15.31	4.95	4.07	2.18
SHAMMA1001978	13.78	19.88	28.28	16.09	13.28	16.86	13.57	12.24	23.18
SHAMMA1001992	6	4.83	12.15	2.11	5.12	11.04	3.79	2.55	5.99
SHAMMA1001994	4.27	7.78	10.55	16.33	8.89	5.41	3.19	4.78	5.52
SHAMMA1002008	2.82	3.39	13.27	3.37	3.98	2.92	2.97	2.85	1.68
SHAMMA1002009	3.5	5.4	13.89	4.02	6.32	3.6	2.53	1.71	1.09
SHAMMA1002011	3.21	3.57	5.26	2.05	5.5	3.92	2.9	2.56	2.89
SHAMMA1002022	23.41	32.84	43.13	18.55	24.01	20.69	18.94	17.16	21.03
SHAMMA1002024	4.59	8.48	7.61	4.82	3.7	4.18	3.88	2.33	4.09
SHAMMA1002032	22.72	24.3	34.55	17.1	16.95	18.48	20.89	15.73	25.86
SHAMMA1002033	5.27	4.59	9.29	4.37	4.11	4.75	8.94	7.94	5.81
SHAMMA1002041	8.82	11.13	18.18	4.7	5.73	4.73	2.24	3.33	5.35
SHAMMA1002042	4.21	5.9	12.97	1.93	6.7	3.65	5.79	7.29	5.48
SHAMMA1002045	3.58	4.88	12.3	4.27	5.48	4.41	3.9	4.78	2.87
SHAMMA1002047	6.8	7.34	9.55	7.31	12.88	11.48	8.75	3.44	4.92
SHAMMA1002056	2.57	2.86	8.49	3.74	3.21	3.7	1.38	1.89	2.78
SHAMMA1002058	12.8	12.32	19.01	10.51	10.24	9.8	10.2	7.07	10.8
SHAMMA1002060	3.44	2.88	10.21	3.91	3.38	3.73	4.39	1.38	3.75
SHAMMA1002065	5.77	6.94	12.55	3.81	3.34	3.54	3.78	2.53	2.86
SHAMMA1002068	5.43	5.28	11.94	2.41	4.51	4.09	4.21	2.57	4.58
SHAMMA1002070	3.38	5.43	10.65	2.42	2.8	1.57	1.88	0.34	3.17
SHAMMA1002078	3.03	3.94	12.3	1.63	3.98	2.25	2.32	1.3	2.65
SHAMMA1002080	11.92	16.79	18.82	3	9.03	5.78	7.02	2.85	4.18
SHAMMA1002082	8.32	5.53	12.97	8.82	3.85	5.55	2.81	1.42	4.34
SHAMMA1002084	5.95	5.89	9.82	6.01	4.3	5.5	3.1	3.76	3.18
SHAMMA1002087	4.18	2.51	5.36	3.1	3.55	1.79	2.57	2.24	1.51
SHAMMA1002091	4.77	8.45	9.77	6.57	5.69	2.84	2.82	2.99	2.11
SHAMMA1002093	7.77	60.44	12.78	4.52	8.15	3.8	10.88	4.49	8.61
SHAMMA1002095	2.07	3	10.99	1.29	1.85	0.7	1.89	0.49	1.88
SHAMMA1002108	11.08	15.82	32.13	20.98	14.75	19.73	14.58	13.23	20.21
SHAMMA1002112	11.23	16.95	26.2	18.44	15.29	9.14	18.6	9.46	11.52
SHAMMA1002118	18.81	20.81	31.7	25.23	19.08	12.25	11.43	15.5	24.54
SHAMMA1002119	10	9.39	20.1	14.89	8.54	8.01	7.08	7.05	10.3
SHAMMA1002125	8.81	6.62	15.53	7.95	7.53	5	5.83	5.81	8.41
SHAMMA1002128	28.45	22.62	47.44	8.75	23.38	14.29	28.49	8	13.65
SHAMMA1002128	9.97	18.58	26.22	10.58	12.77	9.47	19.63	7.42	11.63
SHAMMA1002132	6.21	7.25	34.7	5.81	8.67	4.88	8.38	3.29	9.25
SHAMMA1002140	16.11	29.14	22.82	11.88	12.1	5.91	9.18	3.59	4.86
SHAMMA1002142	3.45	5.23	11.28	4.42	6.87	3.75	4.53	0.93	3.45
SHAMMA1002143	16.35	16.97	31.08	20.92	14.17	21.18	9.45	9.06	16.49

46 / 157

【0870】

【表399】

2531

2532

MAHMA1002145	69.23	60.84	77.51	28.82	48.01	14.01	21.49	27.44	27.68	*	**	-	-
MAHMA1002147	5.58	4.1	8.16	3.48	3.17	3.48	3.61	1.58	1.54				
MAHMA1002153	2.82	4.71	7.08	3.28	5.54	4.01	3.33	4.14	1.01				
MAHMA1002155	2.48	3.25	10.88	2.89	3.23	2.81	4.53	1.33	2.05				
MAHMA1002156	1.89	5.61	11.57	0.4	3.84	3.05	2.17	1.45	3.44				
MAHMA1002158	31.08	68.42	55.01	8.28	17.41	13.89	18.74	7.42	4.4	*	*	-	-
MAHMA1002164	1.74	3.48	10.17	3.33	5.39	3.9	5.07	0.22	3.21				
MAHMA1002165	14.98	21.57	31.25	20.08	14.73	17.78	14.07	10.89	17.29				
MAHMA1002170	22.85	29.54	35.22	10.84	9.2	7.67	9.57	17.6	7.88	**	*	-	-
MAHMA1002174	5.8	8.95	10.45	5.64	4.96	7.45	4.49	3.82	4.26		*	-	-
MAHMA1002175	51.5	30.74	138.92	91.4	93.51	53.98	49.84	44.62	29.88				
MAHMA1002180	28.41	35.57	84.21	33.73	28.48	23.51	38.7	27.33	38.33				
MAHMA1002198	37.99	80.63	138.79	18.28	47.65	48.05	13.24	10.75	8.83				
MAHMA1002205	15.14	20.58	52.79	21.27	16.14	13.48	24.47	10.14	16.84				
MAHMA1002206	2.93	3.69	10.31	2.52	3.98	2.18	3.53	0.98	3.24				
MAHMA1002209	183.5	288.42	348.13	282.08	352.1	289.26	189.69	127.24	124.82				
MAHMA1002215	24.38	31.33	40.05	29.39	28.67	24.47	25.62	23.02	20.25				
MAHMA1002219	2.71	0.94	3.73	1.84	2.07	2.08	0.88	0.6	1.35				
MAHMA1002224	15.23	8.24	19.38	12.34	12.12	11.67	6.61	10.38	6.31				
MAHMA1002229	18.48	14.44	27.70	8.51	18.83	11.84	7.32	11.92	8.59				
MAHMA1002230	22	37.1	57.12	14.15	29.28	20.13	22.25	15.48	15.8				
MAHMA1002233	70.38	89.47	122.68	8.19	33.88	39.69	218.52	154.84	147.87	*	*	-	+
MAHMA1002234	7.92	8.85	12.78	4.92	10.39	5.68	11.44	4.89	8.58				
MAHMA1002236	136.07	148.16	220.88	35.9	57.62	87.11	34.48	55.67	55.33	*	*	-	-
MAHMA1002243	20.02	20.33	38.87	38.33	25.12	26.3	21.68	17.83	24.18				
MAHMA1002250	2.68	3.81	6.87	6.79	2.6	4.11	4.54	2.26	2.04				
MAHMA1002253	0.57	1.7	6.1	2.21	1.65	3.88	2.62	1.41	1.88				
MAHMA1002267	9.29	9.08	17.88	10.28	11.78	12.03	8.49	4.34	11.22				
MAHMA1002268	0.57	3.72	6.93	18.18	7.45	4.6	3.59	1.73	3.92				
MAHMA1002269	1.7	2.24	4.35	2.97	8.98	8.51	2.83	1.4	3.35				
MAHMA1002282	2.88	3.78	4.32	2.98	7.03	4.77	3.56	5.53	5.39				
MAHMA1002282	15.43	25.26	45.69	33.39	19.64	22.84	15.48	15.23	21.7				
MAHMA1002283	3.2	3.94	4.4	4.2	3.48	3.24	3.79	4.35	3.43				
MAHMA1002284	9.41	6.68	12.5	3.33	7.99	10.5	5.31	8.44	4.21				
MAHMA1002297	5.35	6.59	11.35	7.95	7.71	8	2.76	2.6	3.09				
MAHMA1002298	3.44	2.01	8.19	4.11	4.61	4.39	2.48	2.98	3.08				
MAHMA1002299	2.99	4.55	14.58	7.08	6.73	6.49	4.69	0.99	4.04				
MAHMA1002308	5.47	9.33	18.02	10.01	8.02	7.61	7.23	3.03	7.9				
MAHMA1002310	9.13	10.94	18.91	13.86	17.88	11.7	14.2	7.14	10.5				
MAHMA1002311	10.67	15.07	18.09	4.83	5.07	8.09	1.95	3.51	2.65	*	**	-	-
MAHMA1002312	30.9	42.08	49.45	34.33	28.68	29.57	22.88	30.71	35.68				
MAHMA1002317	5.72	5.28	10.29	3.9	3.7	4.94	4.31	4.41	4.7				
MAHMA1002319	11.12	14.39	20.39	9.58	9.59	6.6	12.89	13.6	13.85				
MAHMA1002322	2.09	3.24	7.26	1.71	2.86	1.28	1.75	1.09	1.05				
MAHMA1002329	17.33	19.74	32.38	18.64	18.47	17.53	19	10.58	17.8				
MAHMA1002332	2.58	4.23	6.84	4.89	4.12	2.84	2.37	0.43	2.61				
MAHMA1002333	1.45	2.08	7.07	4.55	5.97	6.64	2.39	0.64	2.14				
MAHMA1002335	12.65	18.48	23.72	7.59	6.81	10.8	3.17	6.12	8.64	*	*	-	-
MAHMA1002339	7.58	16	19.05	10.2	7.54	10.35	7.33	10.1	11.23				
MAHMA1002347	10.15	21.39	24.74	8.5	9.44	11.14	10.35	10.5	16.39				
MAHMA1002351	7.61	10.85	21.39	9.12	8.67	7.13	7.44	4.48	10.85				
MAHMA1002352	3.96	6.91	9.29	4.01	3.84	4.53	2.75	1.39	5.81				
MAHMA1002353	8.39	10.78	23.8	6.39	12.46	12.35	7.8	9.12	7.88				
MAHMA1002355	25.93	29.71	53.72	28.58	30.78	23.84	29.82	21.55	27.08				
MAHMA1002356	28.68	42.92	56.16	40.53	46.92	36.89	34.66	27.13	37.63				
MAHMA1002359	9.7	13.9	19.55	15.93	13.24	12.52	6.92	6.02	14.49				
MAHMA1002360	85.31	125.75	125.12	60.2	60.78	78.44	18.48	22.03	30.83	**		-	-
MAHMA1002361	4.68	5.33	13.4	6.92	5.11	6.88	3.35	2.61	7.85				
MAHMA1002362	4.53	5.8	12.93	6.01	7	5.37	5.23	2.68	7.06				
MAHMA1002367	13.67	17.12	24.55	7.42	11.11	9.42	8.07	3.62	6.96	*		-	-
MAHMA1002371	2.82	3.35	9.8	0.89	2.2	2.61	2.45	2.44	1.89				
MAHMA1002380	7.67	13.26	17.64	8.95	9.26	6.58	10	7.71	10.48				
MAHMA1002384	91.44	188.94	87.68	81.83	128.23	76.47	58.65	15.61	10.22	*		-	-
MAHMA1002385	2.04	1.77	7.64	5.82	2.93	3.4	1.44	1.54	1.92				

2533

2534

RAMMA1002390	16.21	20.25	27.48	13.4	13.91	14.48	13.02	8.71	18.23				
RAMMA1002392	4.77	2.83	8.02	7.9	2.39	4.12	3.41	2.08	4.6				
RAMMA1002395	2.99	1.82	9.4	2.48	1.34	2.01	2.08	1.84	0.91				
RAMMA1002399	9.87	15.29	24.4	2.74	8.91	5.25	5.52	3.6	1.73				
RAMMA1002400	72.48	90.25	98.09	48.92	68.69	49.07	35.95	25.31	45.88	*	**	-	-
RAMMA1002409	6.49	9.18	17.47	3.97	6.62	4.36	5.84	1.59	5.43				
RAMMA1002411	9.51	7.57	20.49	9.69	12.54	10.88	11.51	8.91	7.18				
RAMMA1002413	34.97	38.81	65.44	49.38	40.42	34.55	32.31	27.95	41.68				
RAMMA1002417	36.82	49.91	68.59	44.18	31.29	40.31	29.78	24.72	27.25				
RAMMA1002427	6.8	10.05	17.61	10.58	7.31	11.82	13	8.55	8.6				
RAMMA1002428	6.09	6.45	11.72	5.54	5.32	4.74	5.8	2.19	4.39				
RAMMA1002433	14.18	24.47	33.83	12.49	14.56	9.41	13.05	9.04	12.93				
RAMMA1002434	4.26	3.88	13.18	2.12	3.78	2.95	4.67	2.76	3.7				
RAMMA1002445	8.39	8.67	27.55	10.82	15.56	10.5	11.46	8.07	9.95				
RAMMA1002447	4.71	8.42	13.3	4.25	7.94	6.39	4.34	1.12	6.19				
RAMMA1002454	3.91	4.62	7.05	7.12	5.04	6.67	4.56	2.3	3.21				
RAMMA1002481	27.3	34.88	56	30.05	20.73	23.5	29.37	22.7	22.72				
RAMMA1002483	9.24	10.49	15.67	10.5	6.07	8.24	8.27	4.83	7.57				
RAMMA1002464	3.98	2.85	5.66	1.75	3.24	1.37	1.77	2.41	1.96				
RAMMA1002466	4.68	7.93	15.75	8.06	8.89	6.34	8.75	3.85	4.7				
RAMMA1002470	4.68	6.3	14.67	5.95	13.1	4.89	3.64	2.83	2.84				
RAMMA1002475	17.29	22.9	40.72	24.51	29.14	18.55	20.39	12.61	16				
RAMMA1002480	51.48	94.76	100.05	63.97	58.87	37.35	50.56	13.3	33.45				
RAMMA1002485	19.11	15.89	43.85	29.44	26.15	19.25	16.32	13.38	18.32				
RAMMA1002494	22.68	28.77	50.5	24.3	23.02	18.38	23.12	24.83	27.28				
RAMMA1002498	5.27	2.99	4.95	1.99	2.08	1.95	2.18	0.89	1.64	*	*	-	-
RAMMA1002524	11.96	11.69	24.02	7.72	7.78	6.46	10.41	6.02	6.07				
RAMMA1002530	277.32	388.85	679.87	473.31	550.87	394.12	467.12	251.99	328.81				
RAMMA1002538	1.65	4.73	10.72	1.52	2.62	2.66	2.05	0.61	2.49				
RAMMA1002545	2.59	2.37	13.88	1.77	4.03	1.39	1.88	0.09	1.93				
RAMMA1002554	116.83	185.59	278.08	258.19	207.88	207.1	180.88	93.96	83.88				
RAMMA1002556	4.03	6.04	7.54	5.08	3.68	5.28	2.7	3.78	3.11				
RAMMA1002561	15.67	18.37	24.08	5.72	9.69	8.98	6.7	7.18	2.52	*	**	-	-
RAMMA1002565	20.13	19.37	64.78	24.22	20.52	19.19	20.48	14.5	21.19				
RAMMA1002566	5.49	3.28	8.28	2.18	3.2	3.17	4.18	1.97	2.42				
RAMMA1002571	12.37	18.44	38.1	16.95	12.11	12.28	16.65	10	13.05				
RAMMA1002573	4.02	5.04	11.33	1.38	3.48	2.19	3.22	1.15	2.21				
RAMMA1002576	13.78	23.79	37.39	16.06	18.72	15.25	13.9	8.77	10.92				
RAMMA1002584	9.21	12.39	27.02	16.71	13.29	9.15	13.96	6.48	12.24				
RAMMA1002585	6.36	5.36	13.68	11.07	9.12	8.7	7.18	4.4	6.71				
RAMMA1002586	21.75	21.37	47.64	26.29	19.9	20.88	19.88	14.32	18.17				
RAMMA1002589	31.16	38.3	81.02	35.29	38.93	32.17	47.24	33.3	29.17				
RAMMA1002590	8.66	6.16	15.55	9.32	8.84	9.26	5.08	3.24	4.01				
RAMMA1002593	8.47	7.53	30	8.1	8.37	7.45	8.83	8.88	8.47				
RAMMA1002597	11.34	16.94	39.07	13.1	12.78	11.04	12.91	7.38	12.5				
RAMMA1002598	9.23	11.39	28.12	15.11	13.9	10.8	14.04	4.47	10.54				
RAMMA1002603	7.01	8.83	19.81	9.42	12.79	9.59	19.58	3.51	8.05				
RAMMA1002612	0.54	0.54	1.4	4.78	2.04	2.65	2.1	0	0.92				
RAMMA1002617	1.99	2.39	4.19	2.19	2.2	2.69	2.6	1.84	1.34				
RAMMA1002618	16.08	18.63	46.52	25.87	25.59	22.23	19.62	10.7	17.58				
RAMMA1002619	7.82	6.82	18.84	13.61	11.83	9.5	6.72	6.93	5.28				
RAMMA1002622	3.08	2.72	4.49	4.32	2.98	14.83	2.48	0.95	1.91				
RAMMA1002623	2.38	2.14	3.99	2.84	2.39	1.68	1.82	0	1.12				
RAMMA1002625	158.57	183.87	332.03	71.22	106.04	287.72	518.89	291.59	178.62				
RAMMA1002627	6.48	11.48	14.88	11.21	19	9.56	12.93	6.03	9.03				
RAMMA1002628	20.19	19.33	33.94	25.81	18.48	21.46	13.72	14.18	19.09				
RAMMA1002631	4.6	4.73	9.78	3.98	2.11	3.82	1.12	0.35	1.47	*			
RAMMA1002633	64.25	89.28	104.49	122.15	95.78	107.58	88.41	60.3	67.07				
RAMMA1002635	30.48	34.84	49.63	34.43	30.9	28.98	42.01	28.77	46.58				
RAMMA1002637	2.95	5.14	7.08	4.15	3.68	2.57	4.18	1.82	4.8				
RAMMA1002646	3.66	4.18	8.8	3.4	4.56	2.51	2.73	1.37	2.49				
RAMMA1002648	136.47	120.19	147.15	150.69	135.33	106.18	124.23	88.88	146.01				
RAMMA1002650	1.26	1.85	1.15	1.57	0.72	1.64	0.26	0.46	0.54	*			
RAMMA1002652	32.12	44.28	48.88	48.41	32.68	36.9	24.05	25.91	28.22	*			

48 / 157

2535

2536

MAHMA1002655	11.4	14.99	17.9	15.12	12.04	14.83	12.19	8.95	8.63				
MAHMA1002662	8.09	7.03	11.39	9.78	5.9	8.37	9.45	4.05	7.02				
MAHMA1002665	45.5	69.25	80.49	54.25	82.78	58.03	68.39	47.78	71.4				
MAHMA1002671	18.2	21.13	31.24	18.58	22.22	20.8	25.58	15.85	14.12				
MAHMA1002673	11.32	15.65	15.97	11.87	13.24	9.54	13.67	9.79	12.55				
MAHMA1002684	12.93	20.82	16.77	15.69	20.34	11.67	14.21	4.61	5.68				
MAHMA1002685	5.99	5.53	9.67	4.13	4.23	3.82	3.38	2.86	3.46	*	*	-	-
MAHMA1002692	8.9	8.42	9.67	7.25	7.35	7.75	2.28	5.77	1.68	*	*	-	-
MAHMA1002693	9.64	10.93	23.31	11.25	9.77	9.02	7.89	7.85	6.68				
MAHMA1002698	6.28	7.76	12.68	8.95	7.51	8.5	5.99	6.58	6.22				
MAHMA1002699	5.33	9.97	9.52	5.1	18.31	6.85	7.2	5.75	6.88				
MAHMA1002701	26.1	32.61	44.74	24.49	32.98	29.72	38.78	18.9	37.18				
MAHMA1002708	10.78	13.98	19.02	10.41	15.42	12.16	15.37	13.13	12.91				
MAHMA1002711	14.76	23.34	34.41	24.21	29.1	25.59	25.28	14.17	20.72				
MAHMA1002712	11.97	12.21	14.44	7.82	18.08	13.25	6.69	8.26	7.29	**	-		
MAHMA1002716	4.68	7.58	7.57	4.77	4.14	4.42	3.66	3.87	3.59	*	-		
MAHMA1002721	14.39	14.75	20.55	11.28	18.99	13.97	10.38	11.19	14.85				
MAHMA1002723	7.42	10.89	16.39	7.15	8.24	4.7	4.46	4.93	6.1				
MAHMA1002727	5.85	7.13	17.65	5.65	6.92	6.82	4.72	3.49	8.01				
MAHMA1002728	209.39	279.55	292.72	287.21	287.38	218.06	240.28	159.35	257.9				
MAHMA1002742	14.02	17.18	15.88	10.59	10.26	10.18	5.71	6.31	6.17	**	**	-	-
MAHMA1002743	13.89	22.29	19.83	22.11	23.47	15.54	12.39	8.78	3.07				
MAHMA1002744	41.83	52.92	97.53	82.43	49.67	62.65	35.95	33.09	52.81				
MAHMA1002746	1.97	3.26	4.93	1.96	3.41	2.27	2.78	1.33	2.98				
MAHMA1002748	7.49	8.51	12.58	6.49	4.9	6.1	6.53	3.04	7				
MAHMA1002754	14.88	17.76	22.12	11.35	12.08	11.06	11.12	5.72	18.48	*	-		
MAHMA1002758	3.59	1.84	11.4	2.58	2.22	2.31	1.99	1.02	1.03				
MAHMA1002762	132.2	169.47	185.48	173.16	185.47	170.91	152.23	100.87	138.71				
MAHMA1002764	11.39	16.4	28.38	10.78	16.28	13.47	14.81	8.65	13.68				
MAHMA1002765	7.34	7.13	16.58	5.42	10.33	6.06	7.41	3.51	6.05				
MAHMA1002769	18.29	13.85	24.43	5.77	11.7	10.08	4.64	4.85	10.59				
MAHMA1002771	3.01	3.94	5.73	4.17	2.21	3.23	5.9	1.8	2.75				
MAHMA1002775	37.45	33.73	58.09	84.46	48.03	62.49	34.89	23.26	53.34				
MAHMA1002780	5.07	6.15	8.67	4.47	3.43	3.09	4.23	2.8	2.88	*	*	-	-
MAHMA1002782	5.1	4.96	13.58	4.28	5.88	4.79	6.17	2.83	4.75				
MAHMA1002795	4.5	8.79	10.84	2.77	4.7	2.59	3.8	0.29	3.67				
MAHMA1002796	5.94	10.22	11.38	3.51	7	2.99	1.75	1.74	3.7	*	-		
MAHMA1002805	7.95	9.3	18.25	5.36	11.22	10.85	9.08	3.61	6.4				
MAHMA1002806	3.52	2.38	6.89	3.71	4.28	2.38	4.15	0.61	2.13				
MAHMA1002807	18.13	22.17	39.72	27.6	21.23	23.03	24.44	17.51	32.38				
MAHMA1002814	44.71	48.76	77.02	44.68	36.22	39.5	45.03	34.09	60.73				
MAHMA1002817	3.89	4.65	7.4	2.69	1.42	2.37	1.9	1.38	3.31	*	-		
MAHMA1002820	8.85	7.77	12.62	4.45	5.02	3.82	4.58	1.34	5.96	*	*	-	-
MAHMA1002830	214.03	277.57	281.23	229.98	268.12	260.12	258.75	84.59	195.17				
MAHMA1002833	35.45	43.35	56.35	30.19	48.83	31.33	58.07	31.41	58.94				
MAHMA1002835	2.83	4.62	14.15	2.49	4.88	1.91	2.5	0.14	2.48				
MAHMA1002838	11.11	14.42	31.43	18.22	13.58	15.03	12.7	4.89	21.91				
MAHMA1002842	18.2	19.62	33.31	21.85	17.53	15.65	18.5	8.81	13.48				
MAHMA1002843	8.38	8.96	10.43	6.56	6.63	6.17	6.89	4.51	5.73	*	*	-	-
MAHMA1002844	7.03	7.49	9.84	4.13	4.73	3.32	3.78	1.87	4.09	*	*	-	-
MAHMA1002846	2.68	2.31	8.62	2.22	3.45	1.79	1.92	0.92	2.1				
MAHMA1002857	706.2	952.23	1145	1100.2	1170.5	914.39	1224.8	597.31	410.7				
MAHMA1002858	2294.5	2940.8	1870.4	2372.8	1763.7	1187.7	2253.7	1423.1	844.59				
MAHMA1002863	7.34	6.7	14.07	6.73	7.47	6.7	10.22	2.58	6.73				
MAHMA1002868	37.22	42.58	112.8	68.03	44.4	46.77	34.24	23.46	47.24				
MAHMA1002869	27.07	27.29	44.22	74.9	47.13	42.65	24.18	21.4	25.44				
MAHMA1002871	14.38	15.38	24.35	3.88	5.79	4.87	3.41	1.65	2.57	*	**	-	-
MAHMA1002875	10.61	9.93	8.16	4.59	5.33	3.55	2.05	0.93	2.17	**	**	-	-
MAHMA1002879	85.9	153.46	165.38	80.95	121.8	91.59	134.87	73.22	108.93				
MAHMA1002880	3.99	4.47	13.13	3.43	2.59	1.91	2.37	0	2.3				
MAHMA1002881	8.75	7.92	17.49	5.86	8.28	6.29	10.75	4.45	6.2				
MAHMA1002885	3.52	4.77	10.68	3.9	3.4	4.24	2.59	0.83	2.73				
MAHMA1002886	28.73	26.24	71.15	37.73	28.68	31.72	26.26	17.17	36.38				
MAHMA1002887	4.31	6.26	14.8	6.28	3.45	6.36	3.77	1.7	3.28				

49 / 157

2537

RAM1002890	11.31	9.48	18.16	8.58	8.49	18.82	9.15	5.95	8.48
RAM1002892	13.38	11.62	14.89	12.15	11.02	9.05	7.92	7.82	11.8
RAM1002893	8.84	14.35	13.28	7.89	13.97	6.98	10.13	7.23	1.88
RAM1002895	3.34	5.32	14.4	3.48	5.81	1.87	4.28	2.34	3.5
RAM1002898	4.08	5.01	17.31	2.18	5.01	1.78	3.17	0.38	2.02
RAM1002905	4	11.4	20	7.75	8.82	9.71	6.71	1.7	4.48
RAM1002908	2.68	1.92	5.05	4.09	2.8	3.58	1.97	0.64	1.64
RAM1002908	14.93	16.18	26.75	12.12	15.62	14.31	16.05	8.89	18.72
RAM1002909	30.34	34.63	68.19	28.69	18.89	22.65	34.89	20.01	32.96
RAM1002910	13	10.84	27.29	16.01	18.07	12.45	12.7	8.75	10.42
RAM1002925	49.1	23.96	62.42	27.69	36.92	32.28	67.91	43.67	13.09
RAM1002926	25.41	28.93	57.28	38.81	37.53	35.54	28.57	14.19	28.89
RAM1002930	27	34.76	73.68	26.58	25.59	23.42	47.23	25.24	37.21
RAM1002937	65.34	69.27	78.68	108.43	78.87	71.14	57.91	25.1	50.41
RAM1002938	1.84	2.84	7.68	5.03	4.18	4.4	2.05	0.91	2.65
RAM1002941	7.09	8.07	17.41	12.34	8.66	8.04	6.56	3.92	5.68
RAM1002947	16.06	20.72	33.43	32.88	25.81	27	19.39	13.57	20.15
RAM1002984	14.52	21.96	48.68	29.27	22.18	24.68	18.71	13.88	18.48
RAM1002987	6.98	4.19	15.96	5.47	7.15	5.12	3.82	5.07	4.35
RAM1002970	16.09	29.77	65.22	27.81	25.08	19.79	28.98	12.84	29.95
RAM1002971	4.02	4.7	8.36	3.51	4.7	2.27	5.82	2.89	2.48
RAM1002972	3.94	3.63	9.75	6.43	4.91	5.11	4.27	0.9	3.28
RAM1002973	20.88	22.17	31	24.13	18.86	22.18	16.86	11.21	13.94
RAM1002979	129.92	194.32	219.31	161.08	71.59	113.48	55.53	74.31	23.22
RAM1002982	2.18	1.25	7.22	2.29	3.22	3.34	5.83	0.9	3.23
RAM1002987	10.26	12.2	15.88	9.6	9.29	9.6	13.48	8.42	11.93
RAM1003003	10.92	12.48	21.55	9.72	10.82	10.86	12.69	9.52	10.22
RAM1003004	8.35	12	11.41	6.38	11.7	10.34	5.34	4.84	8.08
RAM1003007	5.35	7.51	8.39	5.22	8.88	8.35	8.36	3.78	6.7
RAM1003011	3.33	3.82	5.67	4.14	2.91	4.01	2.44	1.93	2.62
RAM1003013	52.58	50.01	73.47	127.57	69.44	113.38	42.21	37.2	47.45
RAM1003015	4.4	9.22	10.22	6.88	9.17	8.66	8.79	4.74	5.08
RAM1003018	6	3.52	10.72	6.3	3.92	5.12	4.27	1.82	2.44
RAM1003020	9.29	17.63	21.68	11.76	13.24	12.14	9.04	4.99	7.31
RAM1003029	2.19	2.41	7.83	1.47	3.27	2.09	1.98	0.75	2.98
RAM1003031	12.83	17.1	23.08	9.93	17.27	14.35	13.73	7.28	15.8
RAM1003033	7.59	11.58	13.88	8.87	12.75	8.45	8.32	3.68	9.08
RAM1003035	14.44	18.88	35.58	17.37	11.86	13.08	8.41	9.44	10.82
RAM1003039	5.57	8.4	15.11	7.17	7.63	9.67	4.3	4.16	6.95
RAM1003040	22.42	25.75	39.61	22.72	19.78	23.47	30.8	25.19	32.64
RAM1003044	23.58	26.13	43.98	21.92	19.88	21.85	23.01	22.17	23.66
RAM1003047	25.89	30.87	33.12	18.34	23.86	19.64	30.18	18.74	28.8
RAM1003049	2.33	2.17	8.04	1.08	1.14	1.06	1.63	0.08	1.3
RAM1003055	17.22	25.14	35.08	23.45	17.28	15.29	23.26	16.76	22.03
RAM1003058	1.24	3.89	8.92	1.85	1.8	1.77	4.37	0	1.8
RAM1003067	7.01	11.25	14.1	7.31	6.88	7.83	3.88	5.49	8.7
RAM1003068	16.42	21.28	30.29	14.08	18.97	18.54	12.01	11.17	25.5
RAM1003075	6.07	7.88	12.07	6.85	8.1	8.14	7.15	5.28	7.72
RAM1003080	37.69	44.94	70.13	31.56	35.51	31.8	38.3	22.85	38.78
RAM1003082	2.7	6.77	9.29	2.69	2.39	2.86	1.65	0.84	2.49
RAM1003085	11.71	16.38	22.28	5.98	7.13	4.55	3.24	2.25	2.45
RAM1003099	12.83	14.7	25.29	10.89	15.88	9.22	14.89	11.45	12.15
RAM1003102	3.77	6.08	9.09	1.88	4.54	2.6	2.97	0.85	1.92
RAM1003104	2.23	3.23	4.88	5.3	4.9	4.18	1.01	1.61	4.75
RAM1003113	12.38	25.98	27.12	11.65	17.89	10.18	7.19	8.58	13.27
RAM1003126	20.48	25.06	25.6	15.4	13.83	15.96	10.44	8.97	17.98
RAM1003127	11.67	12.89	22.02	11.38	12.69	10.68	9.71	8.8	11.4
RAM1003131	8.9	14.73	17.42	6.88	7.18	8.51	7.06	3.18	4.59
RAM1003135	3.08	2.9	10.48	1.24	2.69	1.3	3	1.87	1.55
RAM1003140	5.48	8.32	15.55	2.89	4.63	3.63	3.24	3.7	5.1
RAM1003145	4.31	8.5	9.12	2.65	6.5	1.58	4.41	3.82	3.28
RAM1003160	4.32	7.47	10.68	5.78	4.99	3.78	3.2	3.41	6.87
RAM1003164	2.93	3.77	8.99	2.37	2.94	2.85	2.88	0.86	2.1
RAM1003155	35.28	33.7	28.41	18.29	14.13	13.99	27.83	12.47	20.55

50 / 157

2538

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2539

2540

RAM1003157	9.8	11.92	16.49	8.35	7.14	7	5.32	5.07	6.53	*	-
RAM1003163	8.69	5.07	11.11	7.07	6.08	5.64	2.87	4.3	4.05	*	-
RAM1003164	9.65	8.42	12.05	4.3	5.95	3.77	2.65	1.18	3.38	*	-
RAM1003166	13.18	17.5	21.98	14.35	15.87	10.73	8.74	5.62	7.88	*	-
RAM1000010	22.17	30.34	35	8.08	28.78	14.32	17.42	5.42	10.61	*	-
RAM1000016	6.11	10.91	22.1	11.04	8.39	6.1	4.87	1.93	11.92	*	-
RAM1000043	10.01	15.51	20.02	13.55	11.04	7.12	6.17	3.99	7.07	*	-
RAM1000045	170.84	205.1	142.2	172.25	138.41	173.74	84.84	88.64	18.58	*	-
RAM1000054	36.72	48.14	61.65	45.82	42.67	38.97	50.28	31.68	29.41	*	-
RAM1000076	7.95	11.11	16	8.53	7.69	7.72	11.69	4.88	8.85	*	-
RAM1000085	4.15	2.88	11.91	2.38	4	2.05	3.15	0.81	2.03	*	-
RAM1000091	7.05	9.84	23.41	7.46	11.28	6.73	9.81	4.19	5.49	*	-
RAM1000018	6.2	11.24	12.18	4.32	8.37	5.02	4.98	1.12	2.49	*	-
RAM1000032	6.17	20.65	14.02	3.2	3.21	6.55	7.1	3.44	6.93	*	-
RAM1000035	5.32	12.32	11.65	5.42	4.73	4.64	3.82	4.31	2.73	*	-
RAM1000037	6.94	9.49	8.81	3.59	2.53	4.32	3.57	2.25	1.49	**	-
RAM1000039	17.35	20.78	16.95	11.31	10.08	7.38	8.83	5.7	8.35	**	-
RAM1000042	285.1	604.24	359.39	418.78	388.15	125.88	306.06	229.38	295.8	**	-
RAM1000055	3.2	5.45	10.27	2.35	2.5	0.71	3.16	1.31	1.79	*	-
RAM1000059	6.89	5.28	12.3	2.6	4.12	1.14	3.74	1.4	2.81	*	-
RAM1000082	3.25	17.72	14.67	2.77	4.02	2.38	6.97	0.86	3.27	*	-
RAM1000085	129.93	148.35	187.39	51.11	120.84	109.9	91.66	120.8	204.01	*	-
RAM1000086	8.15	10.03	11.83	3.29	5.25	5.44	3.51	4.17	3.47	*	-
RAM1000071	683.12	662.77	828.24	110.41	580.78	269.28	351.73	417.5	139.33	*	-
RAM1000080	24.34	23.12	24.37	3.58	11.29	8.21	9.27	5.24	4.03	**	-
RAM1000088	10.65	12.97	11.63	2.83	5.32	3.9	5.84	4.4	2.88	**	-
RAM1000092	20.13	29.95	35.15	38.08	29.43	24.81	19.3	14.65	20.49	**	-
RAM1000118	1.37	1.52	8.04	0	0.87	0	0.38	0	0.79	*	-
RAM1000119	1.84	9.53	13.52	3.77	3.18	2.59	3.82	0	1.09	*	-
RAM1000121	4.53	3.89	4.87	2.15	3.4	5.49	1.08	0.78	1.77	**	-
RAM1000122	12.08	9.02	14.76	4.72	5.56	4.77	3.17	7.39	6.45	*	-
RAM1000127	4.86	3.09	8.85	2.85	2.99	2.48	1.52	1.46	1.57	*	-
RAM1000131	2.61	0.69	3.44	0.91	1.56	0.43	1.25	0.9	1.02	*	-
RAM1000132	14.01	8.88	18.39	14.15	9.4	7.78	17.49	15	5.09	*	-
RAM1000153	5.01	3.78	14.31	1.42	4.08	2.81	5.68	1.59	2.53	*	-
RAM1000184	281.4	361.08	212.97	228.94	161.22	121.63	212.54	317.83	41.9	*	-
RAM1000186	2.47	4.7	12.58	1.83	2.84	2.03	5.08	0.92	2.79	*	-
RAM1000187	13.51	18.82	27.58	4.35	7.1	11.13	3.58	3.22	8.78	*	-
RAM1000199	2.37	2.51	5.33	3.1	3.18	3.27	2.76	0.65	1.97	*	-
RAM1000213	14.38	13.17	27	9.3	8.17	9.81	8.4	3.99	3.83	*	-
RAM1000216	127.65	93.55	105.55	47	125.55	90	93.27	92.32	37.71	*	-
RAM1000218	17.5	12.81	37.21	8.7	7.88	13.12	12.93	9.58	4.51	*	-
RAM1000224	38.8	21.38	99.03	21.88	23.08	30.11	58.22	41.03	10.49	*	-
RAM1000238	128.23	106.27	224.67	38.44	158.46	117.76	233.25	180.75	49.08	*	-
RAM1000242	0.21	0.74	2.56	0	1.12	0	0.56	0	0.78	*	-
RAM1000244	7.07	7.69	14.14	11.09	6.98	9.63	3.01	2.46	3.87	*	-
RAM1000252	3.63	3.88	6.34	5.75	3.24	4.11	5.02	2.57	2.44	*	-
RAM1000256	13.89	13.11	17.88	12.49	12.18	16.55	9.96	9.49	3.6	*	-
RAM1000257	7.26	4.79	22.28	12.27	7.82	12.93	3.88	4.5	1.41	*	-
RAM1000260	44.37	50.35	69.11	33.91	47.16	50.17	28.8	25.27	25.69	*	-
RAM1000289	4.48	3.88	6.5	3.35	3.23	4.53	2.09	1.54	1.92	*	-
RAM1000271	2.07	1.72	5.77	1.75	2.08	0.52	0.83	0.09	0.38	*	-
RAM1000272	93.8	125.17	184.07	85.39	188.38	110.41	166.71	87.77	97.83	*	-
RAM1000273	44.49	55.95	63.71	32.07	31.24	23.68	15.32	25.55	33.94	*	-
RAM1000274	455.63	719.55	944.44	918.81	390.33	402.82	570.68	748.39	848.53	*	-
RAM1000280	6.37	12.22	16.78	15.15	6.91	15.12	10.38	3.5	4.59	*	-
RAM1000295	0.98	1.35	3.23	1.48	0.9	2.46	0.9	0.93	3.82	*	-
RAM1000300	6.74	6.95	12.1	10.81	8.44	10.7	5.89	3.63	7.13	*	-
RAM1000304	737.06	1023.4	716.47	1038.7	1097.8	489.75	946.67	780.24	888.74	*	-
RAM1000314	4.57	4.01	6.01	3.25	3.98	4.03	4.82	1.32	3.58	*	-
RAM1000318	421.37	294.99	289.4	315.7	574.37	386.7	448.45	209.13	302.89	*	-
RAM1000335	11.86	11.64	18.91	8.84	8.09	12	6.06	5.29	8.48	*	-
RAM1000341	1.84	1.87	2.31	1.53	0.88	0.82	1.84	0.39	2.21	*	-
RAM1000350	8.91	10.88	13.15	11.12	8.93	9.83	7.57	4.14	7.11	*	-

51 / 157

2541

2542

NT2RM1000354	0.4	1.68	2.45	0.51	1.02	0.95	0.87	0.21	0.21
NT2RM1000355	18.03	21.12	26.82	24.57	24.38	11.58	16.5	8.01	13.65
NT2RM1000361	2.57	1.81	2.89	1.57	3.04	3.2	2.49	0.23	2.67
NT2RM1000365	0.53	2.31	3.79	0	3.39	1.9	2.3	0	1
NT2RM1000372	55.88	63.08	56.77	80.59	80.54	75.31	115.69	66.87	19.91
NT2RM1000377	28.22	32.05	43.33	48.89	24.42	20.81	18.03	18.81	14.45
NT2RM1000388	3.72	3.27	4.82	4.15	1.73	3.91	2.28	2.59	2.31
NT2RM1000394	1.58	1.31	3.56	1.15	0.93	0.79	1.9	2.42	1.43
NT2RM1000399	5.42	3.63	8.67	4.2	3.61	3.12	5.84	2.24	2.92
NT2RM1000407	1.68	1.87	7.64	2.01	1.44	1.69	2.06	1.13	3.04
NT2RM1000421	0.74	0.94	3.14	1.35	1.14	2	1.81	0	0.06
NT2RM1000422	134.24	176.38	156.33	143.97	232.18	154.32	116.71	110.46	75.99
NT2RM1000430	4.06	4.21	6.87	5.11	7.78	3.52	6.49	1.64	1.64
NT2RM1000482	17.49	21.49	23.94	21.57	15.84	20.24	13.6	12.18	18.48
NT2RM1000499	11.77	14.5	12.68	4.08	5.8	6.16	2.2	2.63	4.58
NT2RM1000512	60.79	65.64	89.54	15.65	74.29	59.09	19.69	52.23	58.59
NT2RM1000519	108.64	108.15	106.18	60.23	134.25	94.69	58.45	58.56	60.86
NT2RM1000527	51.37	48.34	68.1	100.88	84	85.38	54.81	19.74	44.25
NT2RM1000539	19.83	27.84	33.74	35.02	28.58	31.96	18.52	8.8	16.69
NT2RM1000542	2.91	2.41	11.46	0.76	1.72	0.85	1.68	0.17	0.24
NT2RM1000553	49.27	35.57	83.05	73.97	60.28	58.45	40.5	24.9	19.69
NT2RM1000555	85.13	121.39	124.03	87.28	76.76	74.83	41.33	57.49	77.7
NT2RM1000558	26.75	25.54	23.49	38.55	31.73	38.63	21.2	19.34	35.09
NT2RM1000563	9.05	10.86	14.09	10.89	8.42	9.24	7.48	6.52	10.32
NT2RM1000566	6.73	8.38	16.32	6.55	8.24	8.4	8.98	6.08	8.24
NT2RM1000570	125.7	201.18	105.25	41.82	185.51	106.3	185	132.66	153.14
NT2RM1000571	50.08	74.69	65.76	14.46	84.43	39.98	43.45	24.14	6.89
NT2RM1000574	2.53	1.65	9.83	1.89	3.52	1.88	1.99	0.19	1.58
NT2RM1000580	7.86	7.48	8.12	4.52	12.24	4.74	8.38	3.22	4.57
NT2RM1000620	22.3	25.58	42.74	35.73	32.55	28.94	25.27	21.67	27.34
NT2RM1000623	1.13	1.82	2.61	0.54	0.82	1.17	0.35	0.18	1.57
NT2RM1000620	3.11	2.78	5.69	2.21	1.76	2.12	2.49	0.75	1.9
NT2RM1000633	91.32	89.94	117.87	58.97	108.85	76.85	88.98	107.06	61.97
NT2RM1000634	11.16	13.34	24.68	9.79	14.73	9.21	12.08	6.93	12.7
NT2RM1000642	12.72	15.83	20.12	6.07	8.95	6.42	3.63	2.38	4.88
NT2RM1000647	58.32	165.39	134.62	87.85	112.08	61.29	71.98	52.24	87.23
NT2RM1000648	7.26	13.24	8.89	5.57	12.88	6.5	14.52	3.49	7.15
NT2RM1000650	6.82	9.44	9.48	2.83	6.47	8.87	3.89	5.63	4.1
NT2RM1000681	15.68	18.28	28.1	21.53	19.01	24.34	8.08	7.55	8.54
NT2RM1000685	1.96	1.18	6.92	1.88	1.52	1.45	0.88	0.27	0.92
NT2RM1000689	17.28	18.78	20.82	4.09	4.81	3.69	2.13	1.18	2.17
NT2RM1000672	13.67	18.24	42.69	19.37	17.89	17.56	7.45	2.82	5.99
NT2RM1000681	9.08	18.58	18.31	9.41	17.68	12.82	13.31	6.99	8.03
NT2RM1000691	4.68	7.55	17.83	4	7.1	7.03	5.99	3.47	5.58
NT2RM1000698	12.42	13.98	20.53	4.92	32.5	5.23	15.09	3.16	6.38
NT2RM1000698	5.67	7.09	9.09	7.34	5.78	7.37	2.45	1.86	3.56
NT2RM1000722	25.96	30.69	39.69	30.16	23.23	32.77	24.6	18.35	22.21
NT2RM1000733	32.53	88.64	63.23	38.64	58.43	43.04	42.71	60.17	17.73
NT2RM1000734	43.17	84	45.95	47.88	42.53	19.03	39.69	25.24	10.87
NT2RM1000726	5.67	6.22	17.4	3.29	4.23	3.21	4.45	0.93	1.78
NT2RM1000731	21.17	21.67	40.88	23.29	32.82	14.51	24.97	11.69	10.31
NT2RM1000741	3.88	5.18	10.83	4.4	7.39	2.14	8.17	0	1.39
NT2RM1000742	28.68	28.51	38.7	38.76	29.44	32.74	34.48	38.03	27.02
NT2RM1000744	18.57	21.62	27.17	23.44	20.38	17.29	20.61	18.9	23.17
NT2RM1000745	7.85	8.11	11.72	4.17	4.62	5.27	5.53	3.63	1.14
NT2RM1000747	30.16	38.37	85.95	28.71	33.95	34.03	30.94	18.08	23.76
NT2RM1000752	6.43	11.58	14.15	3.61	7.68	5.78	4.96	2.69	4.73
NT2RM1000767	82.05	87.93	86.38	98.35	104.39	73.49	112.1	58.82	78.42
NT2RM1000770	9.08	11.59	19.18	5.23	6.92	2.95	3.61	2.19	4.43
NT2RM1000772	0.81	1.85	7.5	1.21	3.6	1.85	2.53	0	2.7
NT2RM1000778	107.88	131.01	176.51	63.81	94.24	109.72	76.84	69.35	114.65
NT2RM1000780	6.52	7.85	9.21	2.66	6.05	5.1	2.74	1.38	1.89
NT2RM1000781	6.79	4.22	8.28	1.5	2.8	0.81	2.07	1.28	0.75
NT2RM1000789	63.83	77.27	95.07	93.48	102.47	101.3	60.5	47.17	81.09

52 / 157

2543

NT2R01000820	8.8	11.31	17.21	5.12	7.9	8.45	8.62	2.19	3.3	0	-
NT2R01000822	37.09	50.38	74.58	38.6	34.84	28.86	14.15	4.58	9.84	0	-
NT2R01000811	12.36	9.88	24.62	8.71	10.02	9.68	14.12	5.95	8.7	0	-
NT2R01000828	30.88	47.18	81.3	44.32	50.77	35.09	43.27	14.54	18.43	0	-
NT2R01000829	19.85	28.15	42.48	15.48	18.32	22.11	18.6	10.73	22.04	0	-
NT2R01000831	641.62	837.04	918.82	275.83	388.75	407.08	509.83	541.98	267.92	0	-
NT2R01000832	131.01	121.34	243.58	117.55	172.48	143.88	179.55	110.44	138.22	0	-
NT2R01000834	28.33	24.31	2.89	0	0.85	2.02	26.88	20.39	19.97	0	-
NT2R01000841	24.25	48.24	73.12	11.51	14.12	14.6	18.59	18.34	8.01	0	-
NT2R01000848	14.13	26.98	37.82	13	21.43	18.89	20.71	11.34	13.28	0	-
NT2R01000850	24.68	26.78	38.64	21.16	22.91	25.01	46.02	20.2	29.9	0	-
NT2R01000852	13.97	15.95	23.78	18.75	27.84	15.23	31.21	18.1	21.48	0	-
NT2R01000853	4.2	3.72	6.68	7.37	4.63	5.05	3.89	2.23	2.94	0	-
NT2R01000855	68.98	71.69	82.2	38.85	37.1	51.79	74.88	42.55	21.65	0	-
NT2R01000857	18.68	28.42	44.69	23.2	21.69	23.12	20.54	14.01	19.44	0	-
NT2R01000858	19.44	18.6	24.07	17.38	13.28	16.79	12.93	7.9	10.41	0	-
NT2R01000857	47.06	39.97	81.63	51.39	22.87	30.63	24.25	41.68	28.02	0	-
NT2R01000874	15.63	21.9	42.18	11.95	15.55	12.17	9.14	4.05	6.39	0	-
NT2R01000882	8.95	14.01	25.71	7.66	9.72	9.63	17.08	6.72	11.82	0	-
NT2R01000883	14.26	11.55	31.41	10.17	21.02	6.37	28.25	10.83	26.46	0	-
NT2R01000885	24.96	25.3	31.65	8.98	10.57	16.48	9.38	11.69	11.69	0	-
NT2R01000892	28.28	40.4	46.33	35.83	16.57	27.75	26.99	25.86	21.62	0	-
NT2R01000894	11.18	14.78	12.12	9.67	6.22	9.03	7.82	4.24	5.7	0	-
NT2R01000896	18.49	18.47	21.09	20.98	16.98	15.23	9.37	6.69	9.28	0	-
NT2R01000898	5.15	4.63	7.57	5.14	7.28	6.6	4.73	2.38	2.78	0	-
NT2R01000905	204.63	321.81	294.85	264.7	213.45	208.71	204.71	142.52	180.78	0	-
NT2R01000910	73.69	67.82	69.43	87.95	125.51	72.31	61.51	48.51	44.37	0	-
NT2R01000914	51.65	67.13	48.73	46.18	65.8	39.43	58.15	36.24	73.81	0	-
NT2R01000919	31.63	34.55	49.42	11.97	21.16	18.78	18.17	26.42	20.42	0	-
NT2R01000921	8.87	9.33	11.95	7.94	8.88	8.21	8.17	4.59	6.66	0	-
NT2R01000922	13.29	8.81	21.03	4.57	14.42	11.88	8.29	7.63	6.27	0	-
NT2R01000924	4.71	2.79	7.57	4.61	10.85	2.89	2.79	1.75	0.64	0	-
NT2R01000927	8.24	8.21	11.88	5.08	6.28	6.3	9.48	6.11	7.78	0	-
NT2R01000951	43.38	31.87	51.83	47.29	42.21	49.1	23.28	12.15	26.07	0	-
NT2R01000956	14.05	17.39	13.26	12.15	10.17	18.5	7.97	3.02	8.22	0	-
NT2R01000960	27.19	36.09	41.69	49.18	53.66	39.11	40.71	18.46	20.8	0	-
NT2R01000961	31.7	37.01	55	10.53	21.75	21.52	13.1	26.97	26.19	0	-
NT2R01000982	27.85	38.69	33.9	11.13	12.72	11.99	4.23	8.24	8.72	0	-
NT2R01000973	43.5	24.74	82.26	40.71	70.49	56.15	24.22	34.19	11.88	0	-
NT2R01000978	0.68	0.64	4.81	0	0.42	0.84	0.61	2.08	0.63	0	-
NT2R01000982	16.77	17.99	20.27	10.68	18.58	11.05	7.48	6.36	6.64	0	-
NT2R01000981	7.55	12.77	12.19	10.81	11.04	11.17	5.79	6.02	8.13	0	-
NT2R01000994	39.31	47	23.95	9.37	52.07	40.12	12.98	9.18	11.27	0	-
NT2R01001002	32.74	42.78	50.1	43.61	48.24	33.17	37.45	15.22	13.9	0	-
NT2R01001003	22.14	23.85	35.88	18.55	19.31	27.31	8.71	13.76	14.64	0	-
NT2R01001008	2.98	2.17	4.26	1.07	1.49	2.65	1.82	2.32	1.41	0	-
NT2R01001011	24.49	19.42	39.5	9.16	18.59	21.83	7.1	5.35	13.23	0	-
NT2R01001013	7.68	6.4	15.54	3.26	7.12	6.38	8.21	10.22	6.35	0	-
NT2R01001017	2.88	2.36	7.63	0.68	1.85	2.38	1.32	0	2.89	0	-
NT2R01001018	389.39	678.69	541.38	131	345.59	268.19	208.53	225.12	141.21	0	-
NT2R01001026	7.63	12.1	18.29	9.07	8.15	5.48	3.88	1.52	2.01	0	-
NT2R01001028	4.9	7.17	9.41	5.47	12.08	5.7	11.7	4.88	5.5	0	-
NT2R01001043	26.98	28.41	30.47	32.28	26.93	18.1	7.33	8.19	17.19	0	-
NT2R01001044	13.61	15.79	12.73	5.11	11.55	8.29	6.83	6	8.45	0	-
NT2R01001059	17.82	17.16	21.39	29.41	18.53	17.53	21.69	10.63	22.06	0	-
NT2R01001083	4.42	2.79	10.14	1.51	2.68	1.95	3.55	2.61	2.37	0	-
NT2R01001086	5.44	6.95	11.22	3.97	4.37	2.62	5.3	2.25	4.68	0	-
NT2R01001072	2.1	2.09	8.94	0.59	2.67	1.88	2.14	0.14	1.84	0	-
NT2R01001074	4.98	10.47	11.14	5.42	8.47	3.23	4.47	1.73	1.4	0	-
NT2R01001076	7.28	10.41	10.61	10.91	16.92	6.28	16.73	6.71	8.48	0	-
NT2R01001082	15.78	21.18	26.04	16.26	13.72	10.05	11.58	9.75	14.58	0	-
NT2R01001085	1.82	1.24	5.39	0.21	0.47	1.19	1.3	0.65	1.17	0	-
NT2R01001092	38.97	66.81	68.39	64.52	49.81	49.24	59.78	48.35	50.33	0	-
NT2R01001102	10.81	9.79	17.74	9.01	8	8.42	8.99	8.28	5.92	0	-

53 / 157

2544

2545

2546

NT2RM1001101	12.11	14.84	19.71	7.84	12.92	9.54	14.04	7.04	13.98				
NT2RM1001105	4.22	7.99	12.4	1.52	4.88	3.38	1.81	1.94	3.34				
NT2RM1001112	5.28	6.44	10.76	4.28	9.37	3.62	5.38	1.59	3.48				
NT2RM1001116	7.42	8.2	16.09	7.99	9.99	7.83	8.82	3.88	3.94				
NT2RM1001122	13.95	16.42	25.8	14.85	15.41	13.67	7.73	7.37	13.09				
NT2RM1001136	4.38	9.14	8.72	2.18	2.27	3.09	2.35	0.89	0.34	*	*	-	-
NT2RM1001139	10.44	18.1	18.58	8.95	9.3	8.97	8.52	6.5	7.34	*	*	-	-
NT2RM2000000	3.68	2.81	8.38	4.11	1.99	1.48	3.39	1.08	1.78				
NT2RM2000010	19.98	28.87	32.42	17.38	19.88	13.54	34.18	15.87	15.17				
NT2RM2000013	13.11	32.27	43.35	20.14	26.85	16.98	10.37	4.84	2.67				
NT2RM2000030	111.85	65.82	177.02	160.21	118.48	166.68	73.81	57	69.05				
NT2RM2000032	47.5	43.12	78.78	78.26	63.66	52.78	55.62	28.2	43.38				
NT2RM2000039	27.93	34.32	46.51	34.09	25.35	35.27	22.38	22.42	18.15				
NT2RM2000042	16.44	18.82	11.91	10.78	6.19	11.58	11	10.28	4.44				
NT2RM2000092	22.91	29.95	35.12	39.39	23.39	35.68	21.42	15.87	21.82				
NT2RM2000093	3.68	4.4	7.65	2.5	2.81	1.79	2.27	5.17	2.19				
NT2RM2000093	9.7	29.83	14.41	14.47	18.41	7.29	12.45	10.38	11.18				
NT2RM2000101	92.12	98.68	118.62	114.81	128.83	118.95	69.62	62.13	78.59	*	*	-	-
NT2RM2000104	82.95	69.41	129.33	110.95	90.64	74.09	72.52	52.94	63.78				
NT2RM2000124	8.36	13.36	22.64	10.6	20.3	10.85	19.19	7.38	16.25				
NT2RM2000155	8.81	18.1	14.45	7.85	7.12	5.48	4.06	1.85	4.16	*	*	-	-
NT2RM2000191	45.98	51.15	77.89	79.82	71.81	59.62	35.22	25.83	44.53				
NT2RM2000192	4.72	2.97	6.83	2.91	2.72	2.96	2.76	1.81	1.47				
NT2RM2000239	13.1	14.91	17.38	3.56	5.21	3.85	9.92	3.38	2.39	**	*	-	-
NT2RM2000241	10.79	19.6	19.09	8.89	13.98	6.49	11.48	3.01	8.51				
NT2RM2000250	5.16	6.11	16.92	5.37	6.41	3.58	6.45	1.4	5.69				
NT2RM2000259	7.84	17.03	22.56	4.92	6.83	6.63	8.82	2.03	2.42				
NT2RM2000260	9.47	10.65	8.59	4.97	5.31	6.44	6.43	4.5	4.7	**	**	-	-
NT2RM2000265	1.13	1.27	5.15	0.34	1.13	1.33	1.14	0	0.82				
NT2RM2000287	47.68	55.1	96.28	31.39	38.17	31.93	36.1	33.17	17.11				
NT2RM2000308	68.48	31.19	72.95	23.2	54.18	20.2	25.43	22.47	8.61				
NT2RM2000312	32.44	17.61	49.51	8.48	13.33	15.12	25.48	21.27	10.02				
NT2RM2000322	7.98	6.33	18.54	4.19	5.04	3.2	3.47	0.77	1.48				
NT2RM2000343	85.23	93.08	177.82	122.74	138.14	108.3	83.09	48.31	57.15				
NT2RM2000359	6.22	10.59	23.8	9.56	9.52	5.13	12.03	2.74	6.22				
NT2RM2000362	121.28	140.62	248.7	173.82	223.68	249.97	108	78.11	121.29				
NT2RM2000383	3.84	6.52	9.98	8.57	3.72	4.83	5.89	2.02	2.62				
NT2RM2000388	26.44	18.81	43.02	11.49	11.34	10.48	8.57	3.39	3.24	*	*	-	-
NT2RM2000371	547.25	508.14	752.99	148.13	425.89	273.34	355.28	228.93	135.68	*	*	-	-
NT2RM2000374	12.98	13.63	33.5	14.47	12.05	8.49	18.75	11.88	10.92				
NT2RM2000387	30.78	60.24	91.33	10.11	11.64	24.36	10.29	9.16	3.11	*	*	-	-
NT2RM2000393	10.16	9.5	18.93	3.77	7.23	5.64	6.78	3.44	2.74				
NT2RM2000395	3.42	4.7	7.68	4.26	7.48	3.65	6.98	2.09	2.77				
NT2RM2000402	19.66	22.97	34.41	26.49	10.93	25.33	8.72	3.41	2.72	*	*	-	-
NT2RM2000405	3.7	3.85	7.31	4.5	3.17	3.25	3.78	1.94	0.68				
NT2RM2000407	8.18	7.05	11.03	8.23	7.83	7.22	4.38	3.58	3.42	*	*	-	-
NT2RM2000410	6.69	5.24	14.77	10.81	8.59	8.57	4.92	7	5.39				
NT2RM2000420	12.48	12.68	20.53	7.85	6.52	12.68	7.89	6.33	9.37				
NT2RM2000422	13.42	17.5	14.12	8.88	9.58	15.24	10	8.21	5.9	*	*	-	-
NT2RM2000423	15.54	22.01	50.31	29.73	23.33	19.31	31.02	12.98	24.15				
NT2RM2000452	3.39	4.23	12.53	5.47	7.75	4.42	5.91	1.75	4.67				
NT2RM2000469	3.15	2.51	7.63	7.05	2.6	1.83	2.99	3.88	3.38				
NT2RM2000490	12.63	13.33	17.91	18.49	8.81	12.7	9.83	10.71	10.64				
NT2RM2000497	8.31	3.98	9.54	8.32	2.31	5.44	4.36	6.15	6.13				
NT2RM2000502	7.65	10.01	17.15	15.2	7.83	12.84	18.37	18.29	16.37				
NT2RM2000504	8.62	13.61	12.8	14.35	8.61	11.65	11.39	5.05	12.11				
NT2RM2000514	3.24	4.13	6.05	5.11	2.18	2.21	3.1	2.79	1.68				
NT2RM2000522	0.79	1.87	2.28	6.68	1.51	1.86	2.44	0.58	1.28				
NT2RM2000540	38.37	16.58	24.89	25.09	10.1	17.62	15.11	17.92	16.53				
NT2RM2000565	19.88	7.91	19.23	7.94	6.73	6.21	4.49	7.85	6.72				
NT2RM2000566	25.88	29.95	48.95	40.83	31.02	33.77	42.8	32.01	34.75				
NT2RM2000567	5.01	5.28	9.14	7.2	2.55	2.31	2.41	4.02	2.28				
NT2RM2000568	11.75	12.7	18.59	10.42	10.18	8.54	12.04	11.54	10.63				
NT2RM2000577	23.09	35.16	46.55	20.42	16.81	17.63	9.85	9.61	18.29	*	*	-	-

54/157

2547

2548

NT2002000581	18.64	23.78	26.4	16.81	10.83	8.04	10.45	10.74	12.79	*	-
NT2002000582	34.62	18.91	29.03	32.54	12.67	23.98	20.07	22	22.14	*	-
NT2002000583	208.96	155.13	213.16	202.22	154.41	234.18	169.86	193.18	204.6	*	-
NT2002000589	21.73	20.45	26.61	21.73	16.89	13.34	11.8	15.89	14.79	*	-
NT2002000594	3.88	3.97	9.16	6.31	1.87	1.58	2.08	3.22	2.03	*	-
NT2002000599	23.3	37.43	42.88	41.55	34.51	20.59	19.28	20.01	29.72	*	-
NT2002000609	11.28	11.34	20.18	17.86	7.32	6.53	10.38	8.31	8.85	*	-
NT2002000612	12.78	13.34	16.09	12.28	9.14	7.94	10.91	11.84	11.22	*	-
NT2002000622	141.09	153.93	183.67	159.94	144.1	158.9	166.97	168.17	180.75	*	-
NT2002000623	10.1	4.82	12.61	17.89	6.78	12.88	6.19	8.79	10.68	*	-
NT2002000624	40.85	32.48	58.81	29.49	39.42	24.44	20.09	35.85	21.95	*	-
NT2002000632	4.22	16.93	13.59	10.55	4.72	5.77	3.71	9.75	42.27	*	-
NT2002000635	17.3	14.29	24.16	11.9	11.44	7.82	6.12	9.2	22.13	*	-
NT2002000636	8.64	6.1	10.92	8.1	4.79	4.03	4.23	7.12	5.02	*	-
NT2002000639	12.38	11.39	15.28	10.85	8.59	7.7	5.72	6.03	7.49	**	-
NT2002000649	21.7	22.5	30.17	18.52	14.23	12.19	12.61	12.84	13.29	*	-
NT2002000658	22.82	43.08	46.88	26.73	41.37	32	30.81	19.44	22.2	*	-
NT2002000660	14.51	13.27	29.92	14.04	12.55	18.72	19.19	15.37	13.11	*	-
NT2002000669	10.87	10.69	21.85	10.05	8.68	6.68	4.07	7.21	13.36	*	-
NT2002000680	95.32	133.89	72.27	67.79	75.12	39.22	34.72	103.73	96.38	*	-
NT2002000681	8.29	7.99	12.68	7.17	5.02	1.88	1.67	3.91	2.3	*	-
NT2002000714	22.27	18.86	27.51	17.39	20.26	19.54	14.42	10.68	8.4	*	-
NT2002000718	7.47	11.22	8.78	7.49	4.21	3.74	8.42	4.3	4.39	*	-
NT2002000732	30.31	39.24	31.84	18.98	20.8	19.07	19.55	19.14	20.81	**	-
NT2002000735	9.9	11.63	13.44	9.07	5.38	4.22	4.93	3.88	5.02	*	-
NT2002000740	8.93	6.88	14.03	10.18	2.3	7.44	6.84	7.79	7.93	*	-
NT2002000743	6.68	6.8	14.3	6.91	4.28	3.65	5.08	3.66	3.44	*	-
NT2002000772	27.24	33.91	37.84	22.25	28.4	16.62	5.89	15.75	8.63	**	-
NT2002000773	52.46	53.2	70.82	31.56	38.87	27.02	30.18	38.19	37.45	*	-
NT2002000776	68.74	76.62	73.41	45.06	64.43	20.51	36.77	44.34	50.74	**	-
NT2002000784	15.31	17.83	22.47	11.93	15.98	8.63	6.13	10.15	9.45	*	-
NT2002000795	27.14	19.23	31.28	19.3	19.19	9.44	22.75	21.68	23.6	*	-
NT2002000796	7.18	4.13	6.34	5.73	2.37	1.38	4.64	1.74	1.29	*	-
NT2002000798	220.92	157.62	241.6	267.45	173.39	313.04	164.81	215.06	234.9	*	-
NT2002000801	208.47	181.87	250.83	201.4	210	278.53	117.76	135.99	178.08	*	-
NT2002000821	19.71	21.75	22.88	19.91	8.23	8.65	3.33	5.35	4.94	**	-
NT2002000829	32.72	37.9	42.47	37.81	27.12	23.87	12.82	16.2	13.48	**	-
NT2002000837	12.85	10.7	21.49	8.33	5.59	2.89	4.75	4.19	3.98	*	-
NT2002000924	66.24	48.22	89.49	72.54	62.16	39.92	33.79	29.08	40.4	*	-
NT2002000930	48.55	63.69	52.84	45.58	61.8	35.78	27	25.97	35.7	**	-
NT2002000937	31.8	58.4	47.42	22.54	35.77	16.38	12.6	7.67	4.38	**	-
NT2002000939	6.82	8.82	14.57	7.32	3.05	3.47	2.42	2.17	3.83	*	-
NT2002000942	734.65	670.2	631.32	344.25	467.07	301.67	150.27	478.82	381.55	**	-
NT2002000951	4.26	4.86	13.05	5.55	2.39	1.48	1.13	1.84	1.19	*	-
NT2002000952	9.87	43.75	19.11	12.99	13.49	9.68	6.64	9	6.45	*	-
NT2002000966	187.84	165.71	228.75	214.51	217.85	161.14	133.25	108.26	108.39	*	-
NT2002000973	104.1	257.68	179.14	129.48	180.8	103.68	80.16	78.38	82.31	*	-
NT2002000983	44.05	48.21	49.47	28.35	25.15	19.95	21.95	14.49	14.54	**	-
NT2002000984	14.04	18.71	14.65	9.74	5.64	4.75	5.34	3.16	3.97	**	-
NT2002000994	36.03	29.02	31.03	22.45	13.38	11.64	9.34	22.92	20.85	*	-
NT2002001004	87.18	99.34	120.85	105.48	104.78	118.66	75.52	78.54	83.2	*	-
NT2002001022	485.07	829.81	488.85	279.37	481.13	292.23	411.29	608	337.91	*	-
NT2002001035	118.48	128.84	135.94	89.22	98.88	78.14	85.64	92.65	116	*	-
NT2002001038	39.1	63.09	55.15	35.29	31.08	27.29	33.8	26.97	58.89	*	-
NT2002001043	17.78	17.9	22.19	14.07	14.29	12.96	14.08	13.2	13.27	*	-
NT2002001050	16.38	10.75	11.34	9.74	5.11	3.09	5.76	4.31	3.76	*	-
NT2002001055	7.49	7.2	8.19	5.79	2.69	1.59	3.42	2.92	3.04	*	-
NT2002001065	18.94	19.43	21.19	16.28	10.87	7.49	3.44	9.8	10.97	*	-
NT2002001075	352.65	309.42	355.51	314.53	188.83	298.2	178.81	439.36	452.82	*	-
NT2002001083	7.08	9.84	12.58	9.39	11.2	5.35	6.48	10.07	4.34	*	-
NT2002001100	122.53	127.49	140.57	168.49	157.74	188.39	108.36	105.08	125.08	*	-
NT2002001105	19.7	23.88	18.02	9.87	6.9	7.5	7.31	7.33	7.99	**	-
NT2002001109	20.13	43.33	44.99	10.64	14.91	10.34	19.4	14.81	19.72	*	-
NT2002001110	32.89	28.37	36.47	39.19	23.1	32.21	27.03	23.61	20.38	*	-

55 / 157

2549

2550

NT2RM2001128	10.89	14.18	16.02	9.38	4.33	6.64	7.81	4.58	6.85	*	*	-
NT2RM2001131	35.31	23.37	27.48	39.22	24.25	32.19	17.26	18.93	15.18	*	*	-
NT2RM2001141	23.07	18.72	24.48	15.15	8.42	15.3	18.57	25.01	24.73	*	*	-
NT2RM2001162	7.28	6.34	6.81	7.22	2.67	4.03	3.72	4.85	2.92	**	*	-
NT2RM2001177	18.43	13.33	22.78	9.09	9.55	10.29	11.03	12.82	14.78	*	*	-
NT2RM2001194	18.95	35.2	33.45	15.83	10.75	13.96	9.18	10.24	10.12	*	*	-
NT2RM2001195	22.22	32.77	41.71	19.89	14.02	10.87	14.76	17.07	21.12	*	*	-
NT2RM2001196	12.18	14.33	14.58	8.23	2.8	2.88	2.65	3.74	5.12	**	**	-
NT2RM2001201	112.28	151.54	152.39	113.21	71.67	129.08	182.51	124.84	110.23	*	*	-
NT2RM2001221	15.18	10.19	12.98	12.97	6.39	8.19	4.13	7.92	7.64	*	*	-
NT2RM2001238	7.02	4.76	7.81	7.38	2.47	4.7	2.6	3.8	3.68	*	*	-
NT2RM2001243	28.44	25.47	29.19	20	12.3	9.83	7.01	18.89	8.25	*	**	-
NT2RM2001244	14.84	14.77	26.94	13.43	7.79	12.02	6.48	7.21	5.59	*	*	-
NT2RM2001247	149.17	156.12	167.83	292.88	153.51	292.21	159.18	102.66	137.39	*	*	-
NT2RM2001258	12.98	15.03	17.12	8.24	5.02	5.55	4.85	4.32	8.34	**	**	-
NT2RM2001269	10.95	13.1	12.53	11.07	7.82	4.32	9.04	9.38	5.97	*	*	-
NT2RM2001278	10.38	10.85	11.45	12.81	4.01	9.04	14.51	13.94	18.53	**	+	-
NT2RM2001291	8.24	5.62	12.69	7.3	4.57	5.55	4.22	4.08	3.98	*	*	-
NT2RM2001294	64.54	65.77	111.27	63.88	47.1	58.52	73.02	74.18	85.75	*	*	-
NT2RM2001296	8.24	9.79	14.29	5.81	3.8	3.62	3.42	2.38	2.25	*	*	-
NT2RM2001302	3.38	6.85	11.75	6.43	3.29	2.51	3.71	3.87	1.57	*	*	-
NT2RM2001306	11.41	14.28	14.68	11.73	5.3	7.08	5.43	5.34	3.99	**	*	-
NT2RM2001312	3.55	4.68	12.72	6.28	2.64	2.63	4.81	2.59	2.39	*	*	-
NT2RM2001319	5.88	9.67	14.24	9.42	5.49	4.3	4.35	3.58	2.75	*	*	-
NT2RM2001324	18.95	7.8	13.78	18.03	4.48	12.42	10.21	14.53	14.88	*	*	-
NT2RM2001345	28.65	21.37	24.71	26.89	8	18.65	5.12	14.99	21.21	*	*	-
NT2RM2001360	17.92	12.3	24.15	13.81	8.68	10.87	6.5	8.95	19.99	*	*	-
NT2RM2001370	10.29	7.05	19.68	11.58	5.81	4.38	5.18	4.87	5.67	*	*	-
NT2RM2001391	7.29	4.99	11.62	6.8	2.82	5.45	4.23	5.86	5.98	*	*	-
NT2RM2001393	12.33	6.71	15.2	11.88	7.78	8.57	5.84	9.03	8.99	*	*	-
NT2RM2001420	8.12	9.09	13.52	7.85	4.31	5.05	3.68	4.14	3.54	*	*	-
NT2RM2001423	17.63	16.34	28	15.84	12.17	12	10.08	5.58	5.91	*	*	-
NT2RM2001424	27.85	17.9	24.35	16.51	10.81	12.48	10.74	18.59	18.89	*	*	-
NT2RM2001482	8.67	5.53	15.28	13.75	8.43	10.52	8.21	10.88	22.33	*	*	-
NT2RM2001499	6.83	5.58	14.88	6.11	6.28	4.16	4.43	4.99	9.08	*	*	-
NT2RM2001504	10.92	6.52	14.85	8.85	7.17	4.98	4.36	8.69	8.68	*	*	-
NT2RM2001524	14.94	10.41	10.91	10.19	8.86	8.87	5.54	8.7	4.11	*	*	-
NT2RM2001530	3.04	4.86	7.01	5.84	1.95	3.09	2.61	3.78	1.3	*	*	-
NT2RM2001533	31.05	33.52	48.05	22.65	24.5	20.88	48.4	35.88	44.09	*	*	-
NT2RM2001540	13.31	12.86	17.79	14.39	9.31	10.73	9.68	7.65	7.91	*	*	-
NT2RM2001544	6.59	3.7	12.06	6.73	1.95	5.1	3.34	3.2	4.05	*	*	-
NT2RM2001547	17.98	17.35	32.25	17.05	12.78	13.81	10.07	14.99	28.08	*	*	-
NT2RM2001558	3.53	3.68	12.29	5.32	1.92	2.05	1.78	3.67	4.24	*	*	-
NT2RM2001575	12.53	9.85	20.98	9.37	6.51	6.57	9.69	11.4	8.27	*	*	-
NT2RM2001582	18.68	18.39	26.78	16.81	11.16	8.85	13.11	10.99	19.58	*	*	-
NT2RM2001588	33.5	33.04	38.82	28.73	19.8	15.21	19.42	19.94	22.31	*	**	-
NT2RM2001592	8.08	9.31	10.24	8.08	7.97	5.3	6.26	2.86	5.52	*	*	-
NT2RM2001803	18.68	34.48	37.52	19.83	12.18	19.75	19.12	15.01	10.88	*	*	-
NT2RM2001805	2.46	1.84	10.05	3.88	0.89	9.29	1.27	2.06	1.52	*	*	-
NT2RM2001811	8.28	11.08	21.04	8.21	4.42	4.51	5.24	5.95	7.18	*	*	-
NT2RM2001813	21.23	41.23	53.28	25.42	31.28	14.27	18.42	48.76	35.82	*	*	-
NT2RM2001826	7.78	5	11.59	5.79	3.19	1.08	2.56	3.93	1.82	*	*	-
NT2RM2001832	12.51	10.55	19.25	10.94	8.41	4.21	3.48	2.87	3.67	*	*	-
NT2RM2001833	9.38	12.28	13.2	7.51	9.17	4.14	3.74	4.79	3.89	**	*	-
NT2RM2001835	8.89	10.06	10.54	5.7	4.03	2.66	6.33	4.93	5.79	**	**	-
NT2RM2001838	13.44	12.39	18.22	10.09	11.89	8.86	5.49	5.27	4.75	**	*	-
NT2RM2001837	6.38	4.7	11.19	6.81	1.81	3.12	2.09	2.98	1.51	*	*	-
NT2RM2001839	12.13	12.21	20.77	10.63	8.64	7	4.1	7.5	7.85	*	*	-
NT2RM2001841	8.65	5.25	15	6.44	4.6	3.79	5.03	10.65	9.68	*	*	-
NT2RM2001843	13.92	14.18	22.25	10.74	7	2.7	2.29	4.88	4.18	*	*	-
NT2RM2001848	39.15	45.91	36.48	28.82	23.84	12.46	22.53	23.65	32.08	*	*	-
NT2RM2001852	12.26	10.35	14.74	6.98	5.99	3.53	3.25	2.28	3.08	*	**	-
NT2RM2001859	18.4	10.96	13.81	8.84	9.21	1.42	3.18	2.99	3.04	**	*	-
NT2RM2001890	10.08	9.77	12.95	6.57	5.48	2.88	2.55	2.98	2.8	*	**	-

58 / 157

【0880】

【表409】

2551

2552

NT2R02001684	6.02	5.48	18.95	8.08	4.37	8.84	2.47	6.43	6.78	*	**	-	-
NT2R02001688	15.08	19.03	25.63	5.79	8.13	8.44	2.42	3.93	2.92				
NT2R02001670	5.32	7.34	15.89	8.82	2.18	2.28	5.04	5.37	3.96				
NT2R02001671	8.87	8.88	12.78	9.4	4.31	5.72	3.03	4.34	3.98	**		-	-
NT2R02001675	5.39	3.86	9.7	4.63	1.07	1.88	1.29	1.4	0.98	*		-	-
NT2R02001681	10.15	3.72	7.87	5.68	2.43	1.75	3.17	2.29	1.77				
NT2R02001685	5.99	4.42	4.61	4.4	0.95	1.7	1.78	0.44	1.52	**		-	-
NT2R02001688	7.35	10.19	21.98	10.5	8.43	4.5	8.27	2.89	4.73				
NT2R02001695	21.84	15.61	27.76	16.05	12.1	18.2	12.98	24.27	27.99				
NT2R02001696	11.41	12.78	17.84	10.47	4.47	6.07	3.84	8.22	8.43	*		-	-
NT2R02001698	4.46	6.04	9.55	8.82	3.5	2.88	2.2	3.3	2.08				
NT2R02001699	9.41	7.54	15.28	7.73	5.45	4.44	4.63	5.78	3.45				
NT2R02001700	6.01	3.82	15.92	7.19	8.78	1.93	1.65	2.78	1.4	**	**	-	-
NT2R02001704	15.58	11	13.6	7.65	5.94	5.33	7.22	3.63	5.28	*	**	-	-
NT2R02001705	17.58	13	15.93	12.4	7.58	8.26	9.21	8.02	9.01	*	**	-	-
NT2R02001714	6.57	6.52	7.48	5.63	2.25	2.43	2.36	2.21	2	*	**	-	-
NT2R02001716	8.11	8.76	8.16	8.9	3	2.28	1.76	4.87	3.84	*		-	-
NT2R02001718	5.27	4.14	10.17	8.69	1.77	2.4	3.07	2.63	3.19				
NT2R02001723	7.52	14.9	19.44	7.99	8.79	1.58	2.66	6.02	4.35				
NT2R02001727	13.98	15.14	20.23	10.17	8.39	8.84	18.21	10.41	9.88	*		-	-
NT2R02001730	19.8	22.32	24.32	11.37	7.04	10.08	9.63	10.59	8.89	**	**	-	-
NT2R02001738	24.21	32	34.82	11.52	20.89	8.94	10.25	6.61	7.37	*	**	-	-
NT2R02001743	11.04	12.3	13.73	7.4	4.32	2.84	7.29	5	5.26	**	**	-	-
NT2R02001753	33.97	33.52	45.08	27.22	20.28	23.57	19.89	16.73	21.61	*	*	-	-
NT2R02001756	5.43	4.23	7.05	8	2.04	4.11	2.32	3.16	2.92	*		-	-
NT2R02001760	42.97	48.84	32.99	27.92	15.82	19.33	23.92	44.69	49.12	*		-	-
NT2R02001765	3.08	3.62	8.89	4.79	2.41	2.91	4.69	3.78	2.84				
NT2R02001767	166.3	123.32	123.27	161.08	199.81	185.95	99.49	113.99	129.4	*	+	-	-
NT2R02001768	10.21	7.85	24.02	6.47	5.49	3.96	2.75	3.87	3.14				
NT2R02001771	8.46	12.44	14.25	6.18	3.68	4.08	3.38	4	3.1	*	**	-	-
NT2R02001778	7.68	9.54	13.49	7.21	3	4.06	5.69	4.22	4.13	*		-	-
NT2R02001782	8.45	18.28	12.43	10.97	4.91	8.55	7.19	5.87	4.31				
NT2R02001784	2.33	3.38	2.94	5.28	0.42	1.28	1.88	3.54	2.58				
NT2R02001785	28.54	25.99	26.3	14.79	12.05	14.71	11.42	23.99	14.87	**		-	-
NT2R02001792	9.85	6.48	8.81	6.44	5.88	3.8	3.68	3.95	3.88	*		-	-
NT2R02001795	6.29	3.46	12.45	6.74	1.68	4.68	1.95	2.86	2.03				
NT2R02001797	6.8	5.87	12.89	5.74	3.16	4.45	4.29	3.75	2.07				
NT2R02001800	8.98	9.98	12.02	8.5	3.87	3.21	3.89	2.41	4.68	*	**	-	-
NT2R02001803	9.13	11.59	12.8	14.78	5.7	4.78	5	4.99	8.43	**		-	-
NT2R02001805	6.23	4.93	4.93	5.77	2.81	1.88	2.38	1.02	1.78	**		-	-
NT2R02001806	9.58	8.37	10.47	8.03	2.81	4.18	1.45	2.5	1.81	**		-	-
NT2R02001813	4.64	2.78	4.37	7.6	3.08	5.69	2.05	4.02	2.1				
NT2R02001814	4.12	2.59	2.69	6.32	1.65	3.13	2.04	1.23	1.1	*		-	-
NT2R02001818	2.04	2.24	0.99	4.82	0.99	0.96	0.43	0.73	1.2				
NT2R02001823	3.33	1.51	3.32	5	0.06	3.48	1.75	2.14	2.76				
NT2R02001825	18.28	14.88	18.88	21.81	10.23	18.33	13.27	20.99	16.7				
NT2R02001832	8.66	8.97	9.59	7.39	2.65	5.19	2.03	4.44	3.08	*	**	-	-
NT2R02001839	39.78	58.2	55.7	17.75	23.24	41.75	69.27	61.03	74.92				
NT2R02001840	18.73	9.83	12.67	16.44	10.4	10.18	11.51	19.3	18.18				
NT2R02001851	13.19	13.13	18.5	15.12	5.82	8.1	7.33	10.83	8.67	*		-	-
NT2R02001856	24.09	22.55	30.11	14.9	11.35	11.17	11.7	10.3	11.91	**	**	-	-
NT2R02001867	4.7	5.27	10.74	7.88	2.21	3.72	3.8	4.17	5.58				
NT2R02001868	160.4	263.47	450.85	333.27	292.68	380.02	368.02	285.55	360.35				
NT2R02001879	4.5	3.92	9.65	7	2.4	3.74	3.05	3.17	3.3				
NT2R02001883	0.5	1.23	3.35	4.6	1.17	2.06	2.26	1.81	2.31				
NT2R02001885	5.2	8.33	7.85	8.22	4.29	6.48	4.42	3.65	6				
NT2R02001887	10.27	7.48	19.23	10.21	4.91	6.33	4.19	5.08	5.45				
NT2R02001896	1809.2	1304.9	1708.8	669.09	517.88	708.49	508.04	455.61	977.52	**	*	-	-
NT2R02001902	4.57	4.04	12.43	5.26	2.22	2.22	1.58	3.18	2.3				
NT2R02001903	63.33	63.92	80.39	57.98	54.4	31.38	58.94	50.88	47.6	*		-	-
NT2R02001930	4.16	7.98	10.97	7.26	3.74	5.15	3.87	4.57	4.22				
NT2R02001935	8.31	7.33	13.94	7.14	3.83	4.78	3.89	3.24	2.47				
NT2R02001936	4.31	7.8	13.64	6.57	5.23	5.41	6.78	3.21	3.62				
NT2R02001939	1.41	2.6	6.02	7.8	3.89	3.31	3.08	4.87	3.29				

57 / 157

2553

2554

WT2 R02 001941	10.8	6.4	2.24	11.07	8.67	3.76	4	3.69	5.77	3.42
WT2 R02 001950	10.88	9.24	13.32	9.74	3.68	5.38		5.27	9.54	12.08
WT2 R02 001952	5.22	5.98	18.37	8.77	3.87	2.12		3.56	5.93	8.5
WT2 R02 001976	43.9	27.85	44.8	42.85	38.58	26.62	16.38	23.06	20.88	
WT2 R02 001982	4.3	5.03	8.82	6.09	3.67	2.56		3.61	3.09	3.63
WT2 R02 001983	12.18	15.12	21.04	15.68	19.01	8.03		4.97	6.45	4.73
WT2 R02 001984	20.37	33.54	33.12	14.47	10.58	10.18		8.25	7.68	3.8
WT2 R02 001989	3.88	5.39	6.78	8.08	3.51	2.63		4.78	3.3	2.04
WT2 R02 001996	19.26	18.4	29.01	18.49	15.64	12.47	7.55	13.49	16.15	
WT2 R02 001997	11.71	10.41	18.18	7.7	4.63	5.51		1.14	5.18	8.39
WT2 R02 001998	9.82	8.18	17.59	6.92	5.29	6.42		2.82	8.98	3.65
WT2 R02 001999	18.02	20.28	27.08	10.57	8	5.15		3.81	3.98	3.68
WT2 R02 002003	50.92	53.27	55.93	27	36.02	21.81	18.72	19.18	19.35	
WT2 R02 002004	4.18	6.17	7.32	5.72	2.78	2.44		2.68	3.1	3.84
WT2 R02 002009	12.24	20.57	21.82	12.13	12.38	8.91	7.8	3.92	5.72	
WT2 R02 002014	2.73	6.1	5.88	8.11	3.64	6		6.1	6.78	3.51
WT2 R02 002019	125.13	124.6	184.05	89.22	90.26	64.06	33.48	58.11	68.48	
WT2 R02 002029	13.84	31.19	27.2	28.89	19.23	17.84	11.46	35.39	33.05	
WT2 R02 002030	18.61	18.25	22.9	8.8	11.56	5.38	5.58	12.47	9.07	
WT2 R02 002034	58.72	73.27	60.59	59.82	58.44	27.9	21	20.77	37.34	
WT2 R02 002049	21.26	22	23.61	14.68	15.84	11.74	6.05	9.95	7.32	
WT2 R02 002056	3.27	6.4	6.38	5.47	2.98	2	3.63	3.28	2.55	
WT2 R02 002072	40.82	62.68	39.93	24.05	29.78	18.81	18.47	15.25	18	
WT2 R02 002088	82.61	80.35	97	73.51	51.18	68.26	90.62	55.31	84.87	
WT2 R02 002091	22.22	21.11	34.44	9.78	9.27	5.82	2.71	4.2	5.17	
WT2 R02 002100	12.27	9.78	20.49	8.79	6.55	5.99	3.91	6.96	6.5	
WT2 R02 002109	4.21	5.64	18.28	7.68	3.73	3.18	3.74	2.48	4.42	
WT2 R02 002126	27.51	39.25	34.48	33.04	40.35	13.28	17.62	8.06	21.42	
WT2 R02 002128	4.47	6.67	11.09	4.57	2.71	1.6	2.22	2.89	1.62	
WT2 R02 002129	22.51	19.68	17.11	13.35	16.48	7.78	5.96	8.88	6.92	
WT2 R02 002142	23.72	27.99	22.69	11.44	14.47	7.07	5.7	5.27	5.78	
WT2 R02 002144	5.53	4.42	5.83	7	2.78	3.75	3.22	1.98	3.5	
WT2 R02 002145	12.68	17.57	21.18	8.81	4.88	6.47	2.77	6.22	7.99	
WT2 R02 002153	10.44	8.89	12.47	8.77	5.52	6.62	3.69	3.26	3.47	
WT2 R02 002163	3.34	3.58	10.5	5.65	2.69	1.58	0.8	1.56	1.92	
WT2 R02 002170	9.23	12.24	23.48	10.21	7.83	3.14	3.4	4.81	4.17	
WT2 R02 002178	9.54	6.88	11.83	5.56	4.58	2.92	3.89	2.67	4.48	
WT2 R02 002179	8.2	10.08	9.02	7.04	5.34	2.72	2.79	2.78	2.85	
WT2 R02 002179	5.65	6.68	8.96	6.13	3.31	1.8	2.35	2.99	1.98	
WT2 R02 002179	8.26	8.94	8.28	7.55	3.83	3	2.53	3.14	5.2	
WT2 R02 002179	9.01	16.68	16.49	12.85	7.23	4.54	6.02	8.94	6.77	
WT2 R02 002179	6.62	10.73	17.92	7.67	8.02	5.59	3.83	4.03	4.49	
WT2 R02 002179	12.28	11.64	16.56	11.35	8.87	6.37	4.8	5.99	5.18	
WT2 R02 002179	11.8	12.43	21.01	8.78	6	5.68	4.97	7.35	7.74	
WT2 R02 002179	5.04	4.54	7.81	5.17	3.68	1.33	2.23	2.48	3.79	
WT2 R02 002179	11.8	9.87	12.12	6.06	3.81	1.42	2.55	2.13	2.39	
WT2 R02 002179	5.83	4.84	6.17	8.5	2.45	2.17	3.24	2.41	1.69	
WT2 R02 002179	6.57	3.99	6.62	7.68	2.37	2.81	5.1	3.23	2.52	
WT2 R02 002179	115.68	159.06	153.05	67.65	68	93.3	108.8	133.83	161.04	
WT2 R02 002179	8.81	10.58	19.91	10.24	8.51	6.85	6.71	8.36	8.18	
WT2 R02 002179	12.61	11.63	17.51	7.68	4.03	7.15	6.84	6.41	5.92	
WT2 R02 002179	59.32	60.68	57.77	76.85	24.42	15.38	23.91	11.89	15.71	
WT2 R02 002179	107.02	108.18	125.35	69.13	83.82	75.28	88.4	61.95	98.01	
WT2 R02 002179	21.13	24.18	18.26	10.72	11.76	7.36	11.09	7.92	5.92	
WT2 R02 002179	43.43	55.62	40.03	34.45	40.44	21.34	18.57	11.49	8.97	
WT2 R02 002179	17.23	13.65	15.45	18.01	9.28	15.57	19.85	13.91	16.48	
WT2 R02 002179	10.84	11.13	17.05	9.82	2.18	9.41	7.89	7.53	7.1	
WT2 R02 002179	18.66	20.08	27.47	17.93	14.57	10.2	9.86	18.1	25.3	
WT2 R02 002179	40.87	51.15	69.9	15.35	22.28	15.25	17.48	14.11	15.62	
WT2 R02 002179	50.8	69.26	54.94	17.81	18.82	20.32	24.11	18.36	23.98	
WT2 R02 002179	15.86	23.38	17.16	10.87	6.59	8.39	7.1	5.49	11.21	
WT2 R02 002179	6.96	5.58	6.84	5.68	3.32	1.38	2.92	2.3	3.51	
WT2 R02 002179	63.28	100.72	114.83	36.62	27.49	27.32	87.26	32.69	72.44	
WT2 R02 002179	86.93	107.15	111.07	110.36	46.49	100.89	117.84	46.85	124.45	

58 / 157

2555

2556

WT2R02002764	5.49	12.15	7.39	8.52	3.1	7.78	2.89	4.03	5.78			
WT2R02002772	29.76	33.25	38.82	20.56	12.45	9.21	4.85	18.05	9.09	00	00	-
WT2R02002811	18.39	25	25.21	15	17.51	3.55	6.88	8.61	10.82	0	0	-
WT2R02002818	20.34	18.50	21.22	13.83	9.54	15.73	17.66	20.72	25.81	0	0	-
WT2R02002879	29.35	30.35	32.28	9.31	5.64	8.45	5.48	6.4	8.69	00	00	-
WT2R02002979	45.6	87	85.04	29.24	29.73	11.12	45.71	15.88	60.67	0		-
WT2R02002981	6.39	5.25	5.04	5.12	2.8	1.9	2.21	1.27	2.8	00		-
WT2R02002995	4.58	2.61	3.97	7.89	1.25	3.74	3.35	2.92	4.03			-
WT2R02003031	4.98	5.14	6.7	6.87	2.67	6.19	2.14	4.35	4.33			-
WT2R02003042	32.29	28.58	51.11	28.89	7.79	11.68	9.8	10.82	7.31	0		-
WT2R02003044	5.51	2.76	2.92	6.07	1.88	3.54	2.44	2.91	3.92			-
WT2R02003090	15.61	15.67	25.03	16.18	8.25	19.49	18.74	20.12	15.75			-
WT2R02003095	28.07	26.27	29.38	22.31	14.68	13.94	29.67	24.13	25.38			-
WT2R02003116	19.38	32.85	27.83	12.52	9.52	10.31	8.49	8.73	7.78	0	00	-
WT2R02003222	4.03	3.37	4.48	6.23	1.12	2.42	2.91	3.67	3.87			-
WT2R02003224	23.81	19.9	24.52	18.14	9.42	15.36	24.57	16.25	13.82	0		-
WT2R02003250	119.92	69.2	168.74	153.09	107.43	107.61	63.52	77.81	74.98			-
WT2R02003258	14.27	11.09	23.88	11.45	8.67	5.92	4.04	5.75	7.4			-
WT2R02003262	18.63	6.96	21.6	7.07	3.47	2.7	6.15	5.4	5.15			-
WT2R04000023	18.25	15.39	24.54	14.62	8.88	9.38	9.34	7.69	6.58	0		-
WT2R04000024	5.57	7.1	11.59	9.02	4.21	8.67	3.82	4.08	2.85			-
WT2R04000027	4.73	3.54	9.65	8.85	4.18	2.53	4.65	2.99	2.04			-
WT2R04000030	5.72	8.1	11.8	8.56	5.42	3.94	5.02	4.04	3.02			-
WT2R04000043	5.95	4.77	7.64	13.08	8.98	7.38	10.23	6.96	6.85			-
WT2R04000034	14.03	8.95	17.38	14.71	7.52	11.05	7.66	12.11	11.87			-
WT2R04000046	5.52	6.71	10.27	5.43	2.75	2.16	4.42	4.99	3.5			-
WT2R04000052	3.65	3.61	9.75	4.88	2.31	3.26	1.99	3.02	2.4			-
WT2R04000054	24.72	16.37	22.69	16.92	16.99	12.56	13.57	15.39	11.37	0		-
WT2R04000081	2.45	2.46	7.10	0.32	3.91	3.49	2.41	2.69	0.75			-
WT2R04000074	109.94	111.99	182.79	132.85	134.31	127.91	155.38	114.81	153.05			-
WT2R04000085	16.76	26.39	27.67	18.47	15.63	12.58	12.15	6.78	9.96	0		-
WT2R04000088	5.66	6.66	7.68	8.93	4.39	5.02	8.76	6.10	6.29			-
WT2R04000100	30.87	21.83	35.82	31.94	18.51	25.58	27.68	38.43	48.27			-
WT2R04000101	24.08	19.83	27.89	18.1	9.03	12.08	10.92	15.22	15.58	0	0	-
WT2R04000102	11.73	20.62	40	28.98	36.82	28.28	11.54	34.43	29.82	0		-
WT2R04000104	12.56	10.41	15.15	10.13	9.81	7.37	8.35	10.19	9.76			-
WT2R04000115	5.71	9.33	11.32	11.36	6.79	5.37	7.09	7.13	8.99			-
WT2R04000129	2.55	5.68	5.12	5.02	4.24	3.57	3.26	2.78	3.18			-
WT2R04000139	5.53	8.52	11.67	6.45	6.29	5.11	4.73	4.33	5.31			-
WT2R04000149	3.74	4.09	4.62	8.9	2.02	3.69	4.1	3.49	3.35			-
WT2R04000155	17.15	10.36	17.5	14.41	8.06	9.43	4.52	7.7	8.79	0		-
WT2R04000158	49.23	34.65	52.17	33.08	31.74	35.59	32.83	34.2	32.68			-
WT2R04000187	9.13	4.59	12.54	4.2	2.19	0.82	1.89	2.03	1.71			-
WT2R04000189	116.9	136.33	180.86	211.72	156.35	180.17	146.08	152.04	180.53			-
WT2R04000191	10.04	6.72	12.44	8.1	7.11	6.9	7.89	7.38	7.91			-
WT2R04000197	7.31	6.64	7.63	6.98	3.1	2.95	2.4	2.51	3.17			-
WT2R04000198	13.19	12.13	17.78	14.45	8.75	9.44	12.48	10.4	9.17			-
WT2R04000199	3.72	4.48	6.47	8.34	4.51	5.35	5.27	5.09	6.45			-
WT2R04000201	3.27	5.32	11.9	5.68	2.52	4.69	3.52	3.68	3.02			-
WT2R04000202	7.07	4.86	13.64	7.53	1.78	3.28	4.51	5.82	5.88			-
WT2R04000210	5.45	5.44	15.45	6.67	2.23	1.82	1.86	2.7	2.32			-
WT2R04000215	8.8	9.1	12.68	7.49	4.69	2.59	3.73	4.19	3.48	0	00	-
WT2R04000220	21.3	15.96	19.47	12.48	9.78	7.44	10.31	10.33	6.87	0	00	-
WT2R04000229	4.28	6.1	7.3	4.96	2.64	1.79	2.82	2.28	3.65	0		-
WT2R04000231	12.18	22.17	22.16	12.3	13.79	7.12	11.29	7.48	7.64	0		-
WT2R04000233	21.08	19.41	23.05	29.48	16.53	15.89	23.49	16.8	29.58			-
WT2R04000244	7.04	6.33	12.22	8.72	4.83	3.85	2.95	5.84	4.45			-
WT2R04000251	8.01	7.64	14.53	7	8.84	3.91	3.65	5.44	5.4			-
WT2R04000255	8.99	8.74	14.46	7.34	5.4	4.93	3.56	3.69	2.79	0		-
WT2R04000255	13.77	11.08	20.65	10.54	8.41	4.23	9.38	13.42	10.55			-
WT2R04000283	213.68	220.77	174.14	136.23	205.5	145.83	158.43	132.77	224.43			-
WT2R04000284	85.27	81.74	108.73	78.21	81.89	45.83	89.45	84.2	83.47			-
WT2R04000289	13.22	11	10	10.81	6.15	3.95	7.71	6.59	6.44	0		-
WT2R04000295	9.32	6.67	7.61	11.47	5.8	4.29	6.42	5.34	6.78			-

2557

2558

NT2RM4000306	39.39	31.22	45.46	18.05	17.89	26.02	12.1	14.22	20.93	*	00	-
NT2RM4000307	10.91	9.3	17.74	9.83	6.51	5.04	4.55	5.59	9.36			-
NT2RM4000308	8.25	8.98	13.33	6.03	3.16	3.04	0.89	2.68	1.25	*		-
NT2RM4000311	8.25	9.27	15.68	8.4	4.79	4.48	3.48	4.24	4.11	*		-
NT2RM4000318	7.74	12.3	19.97	11.61	7.91	4.9	8.74	11.58	10.87			-
NT2RM4000324	7.02	4.79	7.97	5.68	2.78	1.11	2.3	1.61	0.69	00		-
NT2RM4000326	9.61	8.92	8.38	6.7	2.41	2.62	3.8	2.41	1.32	00		-
NT2RM4000327	10.62	7.04	10.88	10.24	4.86	9.97	11.64	7.37	9.72			-
NT2RM4000344	28.12	43.85	37.78	23.82	18.35	22.38	18.35	25.83	27.75	*		-
NT2RM4000349	54.86	44.05	74.07	45.08	38.48	49	53.73	62.48	76.26			-
NT2RM4000354	3.83	4.95	10.6	6.31	1.75	1.8	2.77	2.77	2.79			-
NT2RM4000356	5.03	3.65	9.35	5.42	1.79	2.58	1.32	3.38	0.81			-
NT2RM4000366	170.18	124.74	188.28	108.07	122.88	88.88	143.05	122.68	104.4			-
NT2RM4000368	16.94	13.21	17.8	9.95	5.92	5.45	8.44	7.79	10.81	*	*	-
NT2RM4000373	32.64	29.04	37.88	31.16	21.65	24.87	33.11	24.62	25.87			-
NT2RM4000385	6.62	6.28	8.3	8.65	4.71	4.93	6.02	2.91	5.23			-
NT2RM4000395	10.95	13.82	15.83	10.78	7.59	9.02	5.57	7.71	10.22	*		-
NT2RM4000414	4.69	4.37	11.16	5.76	0.91	3.43	2.88	1.6	3.45			-
NT2RM4000417	11.27	10.89	8.64	6.8	4.15	6.41	8.58	1.94	4.82	*		-
NT2RM4000421	10.28	10.99	14.3	8.17	4.31	2.7	2.33	2.02	4.78	*	00	-
NT2RM4000425	91.24	92.57	108.68	64.91	68.82	58.32	76.16	70.13	74.53	00	*	-
NT2RM4000433	6.13	5.12	8.53	7.28	5.88	4.18	2.8	4.18	3.61	*		-
NT2RM4000435	18.9	19.38	27.32	20	14.61	11.77	15.16	14.79	15.69			-
NT2RM4000444	18.5	21.83	23.5	27.68	22.92	21.75	22.28	13.95	20.52			-
NT2RM4000457	13.18	19.12	21.52	10.27	6.06	7.18	7.2	10.64	8.19	*	*	-
NT2RM4000471	11.07	11.03	12.38	8.14	9.49	4.57	3.84	6.85	3.58	00		-
NT2RM4000472	19.66	22.31	17.4	12.44	15.19	12.6	11.64	22.04	27.4	*		-
NT2RM4000485	10.68	18.04	18.73	11.41	9.93	10.39	10.79	12.22	9.89			-
NT2RM4000490	14.19	17.43	15	9.81	4.76	5.97	7.87	5.19	5.16	00	00	-
NT2RM4000496	4.23	8.24	5.23	3.18	0.15	0.82	1.52	0.44	1.12	*	00	-
NT2RM4000505	107.79	139.53	149.56	114.44	75.32	107.4	135.82	107.52	141.57			-
NT2RM4000511	73.01	84.21	82.3	77.22	39.83	68.6	83	54.31	48.28			-
NT2RM4000514	15.58	17.08	20.68	15.1	9.83	18.07	11.41	15.96	14.62			-
NT2RM4000515	15.97	18.43	19.43	7.58	4.12	3.43	3.77	8.97	4.37	00	00	-
NT2RM4000517	351.24	468.1	448.84	338.38	381.29	140.48	205.12	287.92	317.1	*		-
NT2RM4000520	3.58	5.44	6.75	6.2	2.43	4.29	3.08	2.99	1.91			-
NT2RM4000531	7.21	6.94	9.49	7.75	2.88	4.52	5.81	3.99	4.45	*		-
NT2RM4000532	6.65	6.68	4.11	7.27	2.1	1.72	3.22	3.22	3.25	*		-
NT2RM4000533	5.69	4.73	4.92	8.83	1.35	2.87	3.57	2.18	3.55	*		-
NT2RM4000534	4.83	3.29	2.48	5.97	1.61	1.78	3.52	3.71	3.07			-
NT2RM4000583	23.81	15.27	21.44	28.24	15.01	45.53	27.71	31.71	28.82	*		-
NT2RM4000586	9.4	6.85	10.24	13	4.81	13.26	8.88	8.31	8.58			-
NT2RM4000588	24.62	13.96	20.25	24.28	11.25	22.83	7.78	15.36	11.01			-
NT2RM4000589	3.91	1.91	3.09	7.76	1.73	4.77	2.79	4.1	3.39			-
NT2RM4000587	7.59	6.74	8.76	10.89	3.52	5.69	7.38	7.94	10.18			-
NT2RM4000590	2.8	2.23	4.43	8.82	1.35	3.31	2.85	1.87	2.59			-
NT2RM4000593	8.21	5	10.19	8.58	3.29	6.21	6.43	4.7	4.53			-
NT2RM4000595	4.79	4.42	3.36	7.11	6.02	3.43	4.15	2.82	3.87			-
NT2RM4000603	5.4	2.57	9.66	6.3	2.52	2.39	2.28	2.87	2.18			-
NT2RM4000611	28.57	18.91	36.37	25.87	15.39	24.58	19.91	18.95	28.79			-
NT2RM4000616	9.69	7.13	13.2	7.2	3.11	4.07	2.45	3.02	2.58	*		-
NT2RM4000621	3.69	1.04	7.35	6.04	1.62	1.99	3.47	3.26	2.31			-
NT2RM4000648	7.93	6.22	9.62	8.39	3.12	4.35	6.48	5.01	5.06			-
NT2RM4000649	4.41	3.44	7.74	6.28	2.95	1.85	3.85	1.71	2.41			-
NT2RM4000658	5.48	6.14	9.39	8.84	2.67	4.42	4.53	4.69	4.24			-
NT2RM4000681	15.52	9.28	15.4	13.03	7.61	7.72	5.33	7.59	7.97	*		-
NT2RM4000673	13.67	8.93	21.18	9.81	8.09	6.58	4.94	6.04	10.95			-
NT2RM4000674	33.39	14.79	28.45	15.54	18.31	16.48	9.71	12.48	12.12			-
NT2RM4000689	13.32	8.68	20.09	8.6	6.2	5.32	9.33	7.18	6.74			-
NT2RM4000698	3.25	4.45	8.82	7.51	4.85	2.16	4.55	4.72	3.77			-
NT2RM4000700	9.2	12.82	16.94	9.88	10.71	10.68	5.58	6.11	5.58	*		-
NT2RM4000701	37.49	38.63	57.82	37.22	35.87	38.99	50.37	52.92	48.89			-
NT2RM4000712	28.37	34.47	23.93	18.91	14.79	13.59	9.02	8.85	10.91	00	00	-
NT2RM4000717	24.65	10.07	20.07	21.94	7.33	18.21	17.18	23.49	18.21			-

60 / 157

2559

2560

NT2RM4000733	9.25	6.15	14.75	8.98	6.63	6.77	4.25	7.21	11										
NT2RM4000734	6.83	6.23	12.29	7.58	4.56	3.67	3.76	5.75	9.29										
NT2RM4000741	13.34	10.33	15.73	8.84	8.38	7.64	6.59	7.87	3.9										
NT2RM4000744	585.53	451.3	1241	1383.2	1508.5	707.31	777.95	717.55	694.48										
NT2RM4000749	51.3	49.81	57.77	49.48	44.78	42.40	59.41	56.25	54.64										
NT2RM4000751	10.01	10.59	16.82	7.97	7.76	5.05	5.19	5.27	4.64										
NT2RM4000752	14.18	16.62	13.54	10.44	6.56	5.78	8.04	11.07	5.33										
NT2RM4000760	68.65	27.61	44.38	43.74	22.69	40.28	50.15	65.89	60.41										
NT2RM4000761	19.9	12.95	27.34	11.59	8.66	7.67	5.75	12.58	17.08										
NT2RM4000764	28.14	13.4	30.32	14.81	8.8	20.16	18.5	24.89	24.19										
NT2RM4000768	91.29	73.64	83.67	77.4	70.39	93.31	63.42	64.16	68.6										
NT2RM4000778	8.24	6.22	8.88	7.38	4.94	5.13	2.54	2.81	2.48										
NT2RM4000779	13.14	10.85	20.24	8.82	8.68	7.51	2.45	7.69	5.22										
NT2RM4000787	15.56	18.22	17.84	13.21	10.45	11.66	11.77	12.92	11.34										
NT2RM4000790	2620.8	2248.3	2489	2090	3251.6	2752.2	3876.2	1775.7	1268.3										
NT2RM4000795	7.82	4.23	13.52	7.6	3.13	5.92	6.03	8.23	6.4										
NT2RM4000796	26.78	15.24	27.73	18.41	12.57	18.03	14.45	18.92	28.07										
NT2RM4000798	7.01	8.1	18.9	8.2	4	8.33	3.88	7.47	8.07										
NT2RM4000800	5.35	6.72	10.01	6.15	1.63	3.63	2.61	2.89	3.11										
NT2RM4000813	11.25	10	13.48	6.95	4.25	5.74	3.78	4.39	3.13										
NT2RM4000820	28.13	28.55	28.07	22.17	28.99	17.91	22.6	23.35	20.32										
NT2RM4000827	19.03	14.8	13.81	7.34	7.52	5.94	7.29	8.84	3.6										
NT2RM4000830	12.07	12.08	18.71	8.9	8.35	9.39	11.27	7.81	7.47										
NT2RM4000833	11.24	7.17	16.3	8.57	3.24	6.82	2.5	6.31	5.24										
NT2RM4000841	12.98	15.98	14.24	9.21	12.52	7.68	4.07	5.69	5.39										
NT2RM4000846	9.5	7.99	19.32	10.93	5.53	8.29	5.17	10.68	8.48										
NT2RM4000848	7.99	8	13.35	6.98	3.68	2.79	2.7	4.77	2.57										
NT2RM4000852	14.87	14.83	15.83	7.89	7	3.89	3.85	4.67	2.59										
NT2RM4000855	6.4	5.44	7.14	5.82	2.45	2.19	1.77	2.66	1.11										
NT2RM4000859	18.82	15.87	16.78	13.33	8.27	6.7	12.27	10.98	12.51										
NT2RM4000866	19.81	23.16	23.62	14.39	12.92	10.23	4.82	6.2	3.98										
NT2RM4000870	13	13.85	19.75	13.98	14.72	8.68	5.75	8.62	12.65										
NT2RM4000879	41.44	32.18	43.25	14.78	13.02	16.56	3.72	8.51	8.81										
NT2RM4000882	15	11.7	22.42	13.18	7.01	7.9	5.18	5.53	4.92										
NT2RM4000887	41.87	33.25	51.62	28.38	23.39	28.87	18.02	26.87	25.15										
NT2RM4000895	26.8	20.11	33.05	17.45	18.12	7.46	11.88	8.83	10.98										
NT2RM4000897	9.87	8.53	11.07	8.99	4.64	3.44	4.8	3.39	4.04										
NT2RM4000901	10.98	12.11	11.53	7.56	7.1	4.42	5.41	4.23	6.77										
NT2RM4000950	9.78	8.88	10.35	8.13	5.43	3.19	2.51	2.31	1.79										
NT2RM4000965	28.19	32.79	35.21	16.2	12.26	17.85	5.17	10.47	12.18										
NT2RM4000971	97.81	77.57	101.74	132.88	95.27	155.13	87.73	92.85	108.16										
NT2RM4000979	8.96	6.06	14.84	8.87	8.08	6.77	4.88	5.81	4.09										
NT2RM4000987	998.78	831.62	1103.5	625.74	1156.8	778.33	1010.1	1282.3	658.28										
NT2RM4000988	20.87	13	27.71	19.07	13.82	14.2	14.38	15.32	14.84										
NT2RM4000991	41.92	23.13	35.75	37.48	31.48	28.64	21.97	22.75	24.51										
NT2RM4000992	10.32	6.68	8.53	6.07	2.23	2.13	3.02	2.16	2.78										
NT2RM4000996	22.09	29.48	18.01	12.6	11.82	10.41	9.81	3.95	3.92										
NT2RM4000997	28.21	30.93	25.95	19.16	13.92	19.47	16.5	21.82	24.11										
NT2RM4001001	7.63	8.6	15.15	9.35	5.11	5.88	3.98	7.75	5.13										
NT2RM4001002	459.88	441	603.28	484.81	440.37	622.33	521.26	628.75	590.64										
NT2RM4001016	19.09	20.3	24.95	12.05	10.46	13.91	10.42	12.83	14.8										
NT2RM4001025	8.54	7.24	14.2	7.08	5.08	3.26	5.04	3.42	2.59										
NT2RM4001027	4.89	5.86	4.64	4.07	1	1.53	1.18	1	1.23										
NT2RM4001032	13.63	13.78	6.47	5.13	8.41	2.73	4.98	2.77	2.21										
NT2RM4001047	17.2	23.07	17.27	11.83	13.43	8.74	14.05	4.88	2.78										
NT2RM4001049	14.8	12.01	16.43	12.47	5.42	11.65	9.35	13.4	14.12										
NT2RM4001051	23.07	23.96	19.46	18.43	11.09	9.95	3.77	14.4	12.67										
NT2RM4001052	9.72	6.17	11.79	8.32	6.45	2.66	8.57	9.7	6.4										
NT2RM4001053	8.35	8.46	10.82	7.64	6.3	5.31	6.88	7.18	3.68										
NT2RM4001054	15.68	17.03	15.78	11.07	4.7	7.63	9.63	9.71	11.5										
NT2RM4001059	5.24	11.23	8.09	5.5	2.85	2.53	2.48	2.39	2										
NT2RM4001071	23.88	28.79	33.68	18.88	9.28	10.59	17.49	15.69	16.83										
NT2RM4001084	12.62	12.59	13.05	12.4	5.97	9.54	14.16	8.85	15.23										
NT2RM4001092	7.67	7.08	7.41	8.28	5.02	4.82	2.05	6.72	4.2										

61 / 157

2561

2562

NT2RM4001100	10.34	6.87	14.94	8.81	5.31	8.81	7.76	11.53	7.75										
NT2RM4001116	6.46	5.43	9.66	8.08	5.44	5.9	5.27	4.35	5.99										
NT2RM4001119	4.41	2.95	8.51	4.68	4.64	2.33	3.03	2.56	2.24										
NT2RM4001140	6.58	8.81	10.22	5.74	2.39	2.78	3.27	2.64	2.47										
NT2RM4001148	77.43	82.01	87.69	109.01	92.69	80.91	71.63	54.14	60.64										
NT2RM4001151	25.3	23.87	22.4	14.14	12.5	10.51	24	19.61	17.83										
NT2RM4001155	7.7	10.88	9.45	14.26	6.55	9.83	8.68	7.88	7.8										
NT2RM4001167	4.69	2.48	3.77	5.35	1.74	4.87	1.85	2.85	2.34										
NT2RM4001169	10.13	5.87	8.44	9.04	3.06	4.73	3.63	6.09	4.68										
NT2RM4001163	8.69	6.99	8.41	8.85	4.09	5.89	3.67	3.78	3.99										
NT2RM4001187	16.65	10.66	18.41	9.39	4.2	6.99	4.74	5.75	3.81										
NT2RM4001181	8.6	5.61	4.99	6.79	10.14	6.02	8.39	4.7	8.34										
NT2RM4001200	2.58	2.48	3.48	4.38	3.72	1.51	1.91	2.35	0.33										
NT2RM4001203	6.43	5.54	5.53	5.95	1.89	1.97	3.96	4.4	1.73										
NT2RM4001204	9.94	10.43	11.48	7.05	1.89	2.44	3.48	2.67	2.11										
NT2RM4001217	24.89	17.41	30.8	23.78	12.47	10.57	27.41	25.41	29.34										
NT2RM4001245	158.95	84.48	149.16	130.37	118.53	157.67	151.03	141.35	178.48										
NT2RM4001247	4.65	4.91	11.8	4.49	3.6	3.11	6.79	3.17	2.29										
NT2RM4001256	9.61	7.67	12.78	9.69	5.91	6.14	7.73	8.08	8.61										
NT2RM4001250	70.47	64.71	70.62	44.98	48.68	50.08	63.38	45.51	40.88										
NT2RM4001287	5.89	4.4	10.4	6.42	2.33	2.85	4.9	3.94	4.62										
NT2RM4001273	1.34	2.22	3.89	5.02	1.41	1.99	4	2.6	2.16										
NT2RM4001281	8.06	5.61	14.83	10.91	4.33	7.34	4.05	7.62	5.35										
NT2RM4001286	51.19	32.28	36.36	29.23	18.59	18.64	13.09	20.76	31.48										
NT2RM4001290	8.38	5.81	13.36	7.78	4.34	4.65	5.33	4.38	7.29										
NT2RM4001313	98.2	68.64	189.85	90.18	105.33	108.01	85.7	73.37	76.26										
NT2RM4001316	6.16	7.7	10.44	8.55	3.72	5.52	2.76	5.94	3.03										
NT2RM4001320	10.87	13.16	19.79	13.68	8.94	6.89	6.58	5.71	5.93										
NT2RM4001320	20.75	22.22	27.83	18.9	16.02	17.67	25.16	18.45	25.01										
NT2RM4001321	3.2	9.27	5.61	7.64	4.65	3.21	6.77	6.74	7.05										
NT2RM4001325	6.99	10.29	13.71	11.04	4.07	7.04	5.14	7.08	21.4										
NT2RM4001333	11.2	7.57	11.83	8.81	7.22	6.3	2.38	8.09	14.33										
NT2RM4001340	69.12	52.84	78.81	89.02	68.75	87.91	69.71	47.75	38.99										
NT2RM4001344	8.4	4.83	10.13	7.68	4.28	5.37	3.8	5.77	3.72										
NT2RM4001347	414.8	327.84	802.6	388.33	392.69	323.06	332.75	373.98	363.9										
NT2RM4001357	6.26	7.97	8.82	6.78	6.02	4.63	3.22	2.89	2.9										
NT2RM4001360	8.63	8.04	10.78	8.59	6.1	3.6	4.75	7.61	3.39										
NT2RM4001371	15.56	11.85	10	11.21	16.5	13.92	36.9	13.45	6.6										
NT2RM4001377	21.16	11.62	19.22	16.27	7.45	12.65	8.88	10.18	9.45										
NT2RM4001382	18.22	9.22	22.87	11.49	7.37	7.19	3.28	8.65	16.08										
NT2RM4001384	4.53	6.9	14.13	5.13	2.13	2.71	2.62	2.27	2.98										
NT2RM4001400	10.43	8.23	12.46	8.14	6.92	4.37	4.37	6.39	3.72										
NT2RM4001409	4.67	5.55	8.19	6.09	3.26	2.83	3.17	3.27	3.39										
NT2RM4001410	6.69	14.26	12.51	8.64	5.38	3.51	3.98	6.53	3.24										
NT2RM4001411	25.9	30.99	34.08	31.33	32.6	28.51	31.72	28.52	32.83										
NT2RM4001412	625.39	1226.8	1011.2	1004.1	804.98	889.26	1210.2	810.05	943.27										
NT2RM4001414	12.78	8.8	16.28	6.35	2.72	5.16	2.14	4.73	3.09										
NT2RM4001438	63.2	35.98	69.08	66.83	54.08	67.38	42.65	35.42	49.63										
NT2RM4001437	26.18	27.89	39.17	20.31	15.62	18.05	18.35	28.88	22.74										
NT2RM4001444	3.41	3.61	9.9	8.68	3.02	1.89	2.67	2.69	2.26										
NT2RM4001454	19.23	20.25	21.89	16.73	15.4	12.09	15.71	18.45	13.57										
NT2RM4001455	5.88	19	6.35	4.28	2.07	3.65	1.94	1.67	2.98										
NT2RM4001483	8.69	16.03	11.64	7.61	4.41	5.78	3.95	3.81	3.42										
NT2RM4001483	12.59	12.07	8.74	5.56	6.34	2.68	4.94	5.52	3.23										
NT2RM4001495	38.43	24.24	20.4	12.41	18.18	17.63	5.82	16.38	11.58										
NT2RM4001499	22.29	18.15	30.48	16.01	14.05	14.71	4.23	11.08	11.85										
NT2RM4001515	6.81	6.91	14.18	8.91	2.64	4.03	2.9	4.57	2.36										
NT2RM4001519	8.21	4.24	9.85	5.65	1.45	0.78	1.08	3.15	1.34										
NT2RM4001522	8.18	6.53	12.02	8.15	2.62	1.81	2.28	2.64	1.62										
NT2RM4001523	73.84	75.98	71.67	35.81	37.18	20.7	26.91	23.43	33.3										
NT2RM4001550	10.75	6.21	9.98	6.49	6.18	2.2	5.6	2.68	6.03										
NT2RM4001553	11.94	9.8	11.96	9.12	5.14	4.64	3.81	2.93	3.19										
NT2RM4001564	151.88	72	141.69	164.83	103.78	171.98	80.79	76.13	147.24										
NT2RM4001557	61.14	42.32	58.99	56.35	48.65	76.92	34.89	30.81	38.74										

62 / 157

2563

2564

NT2RM4001565	12.7	17.39	26.89	14.14	12.4	12.89	7.45	6.92	4.69	*	-
NT2RM4001566	17.21	11.12	20.03	22.75	14.87	13.43	10.31	12.88	15.08		
NT2RM4001569	3.45	2.79	9.43	4.85	1.24	1.1	2.19	2.38	1.74		
NT2RM4001579	7.75	7.8	9.32	7.49	3.6	1.35	2.94	2.54	2.38	**	-
NT2RM4001582	6.95	5.09	5.63	5.76	1.73	2.34	1.68	1.19	2.1	**	-
NT2RM4001589	118.7	167.05	103.64	41.43	57.6	48.23	27.72	16.73	11.4	*	**
NT2RM4001592	8.17	5.42	14.24	6.52	2.18	4.41	4.13	6.42	4.93		
NT2RM4001594	386.49	312.66	285.69	187	185.32	238.47	180.25	289.69	275.08		
NT2RM4001597	6.23	6.27	13.64	6.79	3.64	4.25	2.98	4.23	3.88		
NT2RM4001605	16.38	18.33	22.25	14.65	11.19	8.87	10.5	12.13	9.23	*	*
NT2RM4001609	23.64	18.9	22.4	13.71	9.55	9.27	6.4	7.54	6.99	*	**
NT2RM4001610	10.55	8.62	11.74	9.35	5.13	4.48	6.11	6	5.63	*	-
NT2RM4001611	14.22	14.7	16.89	10.73	7.27	8.47	6.73	5.19	3.38	**	**
NT2RM4001616	13.74	16.13	14.23	8.24	4.82	5.02	4.84	2.49	3.4	**	**
NT2RM4001622	5.3	5.18	10.27	7.37	3.16	3.88	2.49	3.88	1.85		
NT2RM4001624	7.85	5.37	11.44	5.27	1.81	3.98	3.07	2.88	3.67	*	-
NT2RM4001625	8.28	12.41	10.38	15.36	14.48	2.63	7.15	9.31	10.39		
NT2RM4001628	14.67	21.52	22.9	10.11	13.87	22.31	12.28	12.44	12.55	*	-
NT2RM4001632	325.71	457.93	338.97	242.41	280.06	370.61	594.61	393.87	482.68		
NT2RM4001642	5.97	4.82	5.47	4.64	2.19	2.19	0.94	2.25	0.91	*	**
NT2RM4001647	18.42	12.21	17.82	18.58	14.27	17.66	13.87	14.03	11.49		
NT2RM4001650	27.91	31.97	47.33	38.24	30.14	33.6	22.2	16.78	28.83		
NT2RM4001642	12.58	10.55	10.99	18.39	7.75	15.61	11.38	12.75	8.09		
NT2RM4001646	7.82	11.39	9.77	4.29	5.85	7.48	4.49	4.14	4.38	*	**
NT2RM4001670	25.82	19.71	20.88	17.87	11.22	17.38	18.89	23.01	24.94		
NT2RM4001682	8.89	10.21	10.73	7.71	7.31	3.19	3.94	4.22	3.18	**	-
NT2RM4001710	8	9.89	13.67	6.48	2.86	3.01	1.39	4.38	1.64	*	*
NT2RM4001712	8.13	8.94	9.77	4.06	2.96	2.15	2.17	4.16	1.89	*	*
NT2RM4001714	18.44	20.46	18.31	8.4	3.92	5.08	6.84	3.44	4.9	**	**
NT2RM4001715	22.24	32.3	32.67	17.87	6.68	18.35	15.07	7.8	8.58	*	*
NT2RM4001727	17.83	12.1	16.98	14.9	8.57	19.18	13.54	13.69	14.31		
NT2RM4001731	9.18	5.58	7.2	4.83	1.9	3.44	2.33	4	3.02	*	*
NT2RM4001735	15.03	8.85	13.3	10.72	9.69	6.1	8.95	9.42	9.18		
NT2RM4001739	3.31	4.05	7.1	6.51	1.7	2.71	2.2	2.1	1.76		
NT2RM4001741	5.03	6.39	8.03	4.45	2.85	1.9	3.44	3.9	2.72	*	**
NT2RM4001746	27.18	35.45	35.48	20.34	15.62	12.62	17.33	18.68	21.67	**	*
NT2RM4001754	47.9	62.7	66.48	78.4	39.53	40.99	48.7	37.13	39.58		
NT2RM4001757	4.77	10.74	3.2	6.77	1.73	2.53	2.83	4.79	2.31		
NT2RM4001758	8.89	8.75	6.1	8.32	3.51	6.88	5.98	8.51	8.88		
NT2RM4001768	1.37	0.38	1.12	5.38	1.88	2.21	0.35	1.5	0.9		
NT2RM4001775	6.53	3.93	5.59	8.91	3.97	6.25	4	3.64	3.18		
NT2RM4001775	28.42	28.82	28.25	33.08	15.74	17.3	19.26	20.43	19.52	**	-
NT2RM4001783	4.96	4.55	4.74	5.37	2.6	3.32	2.42	2.8	2.84	**	-
NT2RM4001783	14.95	13.94	19.52	14.23	7.85	12.31	14.48	13.31	17.08		
NT2RM4001810	23.52	24.77	30.47	26.07	19.33	20.61	34.17	38.53	24.95		
NT2RM4001813	9.51	6.59	9.34	11.22	3.7	8.5	12.3	12.03	14.87	*	+
NT2RM4001816	4.65	1.67	8.39	7.01	2.88	5.12	5.55	5.88	5.41		
NT2RM4001819	80.18	35.79	68.43	71.44	40.61	80.22	41.62	48.42	47.06		
NT2RM4001823	10.42	6.03	12.1	10.8	5.46	6.85	5.68	8.09	6.63		
NT2RM4001828	3.67	2.82	6.95	5.88	1.65	1.21	2.42	1.92	2.31		
NT2RM4001835	4.94	4.42	12.73	8.71	1.8	3.96	4.63	4.11	5.11		
NT2RM4001836	5.12	3.81	8.12	6.95	1.63	1.85	2.61	3.45	2.12		
NT2RM4001841	6.73	6.82	10.79	10.33	4.39	6.34	6.7	6.45	7.77		
NT2RM4001842	2.89	4.82	4.7	7.5	2.83	2.4	3.82	3.29	2.69		
NT2RM4001843	48.67	26.12	58.83	48.07	48.73	45.87	28.16	28.11	28.49		
NT2RM4001856	14.1	14.11	19.35	11.87	5.04	7.66	7.47	6.5	8.37	*	*
NT2RM4001858	20.02	12.7	24.11	14.43	7.51	10.18	17.09	15.78	20.13		
NT2RM4001861	14.08	10.48	18.84	13.72	9.53	6.84	12.28	18.05	13.35		
NT2RM4001863	27.15	15.67	30.03	15.24	18.38	17.87	16.63	16.94	11.79		
NT2RM4001865	8.03	9.87	15.29	7.44	8.43	5.34	6.68	5.7	6.85		
NT2RM4001869	12.34	16.35	21.38	13.19	9.35	9.53	19.88	14.54	12.68		
NT2RM4001873	8.61	9.79	10.86	7.15	2.28	3.17	4.21	4.29	2.39	*	-
NT2RM4001876	13.28	18.02	38.33	14.58	13.74	11.65	8.08	13.34	13.31		
NT2RM4001880	22.41	18.46	28.54	18.81	14.09	15.45	13.74	18.99	19.02		

63 / 157

2565

2566

NT2RM4001885	71.31	49.82	81.88	53.98	43.04	45.93	68.83	64.57	48.87										
NT2RM4001889	10.85	7.93	15.01	8.59	4.29	4.67	4.77	6.73	3.99										
NT2RM4001894	30.82	32.08	44.07	33.7	34.28	35.89	33.58	30.42	25.18										
NT2RM4001897	8.76	10.37	11.73	7.79	4.55	4.65	3.35	3.47	2.33	*	**	-	-	-	-	-	-	-	-
NT2RM4001899	4.19	5.18	10.38	7.76	5.5	4.54	4.99	6.17	3.45										
NT2RM4001905	8.04	9.87	9.34	12.13	4.62	8.49	17.28	7.42	9.7										
NT2RM4001922	10.52	7.5	19.46	10.13	4.91	4.77	4.28	7.33	9.75										
NT2RM4001930	42.04	27.93	35.21	26.34	18.3	24.57	35.05	37.8	24.49										
NT2RM4001938	11.88	11.93	17.44	9.19	6.81	6.54	7.09	10.88	7.99	*		-	-	-	-	-	-	-	-
NT2RM4001940	40.15	39.79	42.12	29.48	25.03	28.5	19.62	21.25	18.41	**	**	-	-	-	-	-	-	-	-
NT2RM4001942	7.8	8.46	12.68	8.27	3.74	4.94	3.46	2.55	2.64	*		-	-	-	-	-	-	-	-
NT2RM4001952	2.58	5.71	6.3	5.41	2.95	2.84	2.21	3.1	2.4										
NT2RM4001985	9	9.88	11.74	9.82	12.01	7.33	7.4	10.04	8.64										
NT2RM4001986	61.68	52.74	61.28	70.45	40.89	59.05	68.01	53.06	72.22										
NT2RM4001989	9.53	10.35	15.87	9.41	5.57	8.09	5.17	7.89	12.14										
NT2RM4001974	12.36	21.88	18.69	16.74	8.35	12.5	5.35	17.32	22.51										
NT2RM4001979	23.89	19.24	22.43	14.04	16.93	18.71	17.76	22.89	22.51										
NT2RM4001980	94.16	97.18	94	64.27	69.09	81.77	75.44	72.16	71.83	*	**	-	-	-	-	-	-	-	-
NT2RM4001984	9.87	8.39	11.58	7.27	4.78	4.84	2.35	4.35	2.8	*	**	-	-	-	-	-	-	-	-
NT2RM4001987	7.05	10.61	11.42	6.92	3.16	5.3	2.58	4.02	2.33	*	**	-	-	-	-	-	-	-	-
NT2RM4002013	15.54	16.72	17.68	12.71	8.41	7.52	18.32	18.81	11.68	*		-	-	-	-	-	-	-	-
NT2RM4002018	18.51	18.01	18.38	23.63	14.09	15.67	18.72	13.95	13.04										
NT2RM4002032	8.2	4.97	12.78	7.92	4.07	8.05	2.33	3.53	3.27										
NT2RM4002034	9.06	7.36	15.25	10.05	4.49	7.82	4.17	18.89	9.93										
NT2RM4002044	29.85	20.58	32.58	18.82	16.22	17.19	21.2	22.05	24.1										
NT2RM4002047	20.66	17	25.87	17.48	15.74	9.48	16.1	15.14	16.08										
NT2RM4002054	31.19	30.54	36.58	20.52	17.55	11.13	16.47	15.97	18.8	**	**	-	-	-	-	-	-	-	-
NT2RM4002055	21.78	18.7	21.85	12.08	11.37	8.76	14.04	16.74	17.09	**	*	-	-	-	-	-	-	-	-
NT2RM4002059	10.38	12.77	15.62	6.8	7.63	3.89	4.33	4.04	2.53	*	**	-	-	-	-	-	-	-	-
NT2RM4002061	9	11	9.24	10.93	6.62	5.33	6.85	4.88	4.34	**		-	-	-	-	-	-	-	-
NT2RM4002082	97.19	68.62	99.9	109.04	77.98	163.08	52.62	58.12	71.01										
NT2RM4002083	8.61	5.55	11.75	8.31	3.06	3.85	4.25	4.81	4.84										
NT2RM4002086	7.96	8.56	22.27	12.72	22.89	4.97	2.38	10.48	2.89										
NT2RM4002087	10.67	18.12	21.45	11.17	18.75	7.5	4.23	11.4	5.44										
NT2RM4002073	24.87	29.49	32.35	16.42	15.53	13.85	5.51	8.53	5.15	**	**	-	-	-	-	-	-	-	-
NT2RM4002074	13.13	16	13.48	9.5	9.68	6.27	2.81	4.18	5.74	**	**	-	-	-	-	-	-	-	-
NT2RM4002075	12.68	9.69	13.41	10.12	6.41	6.32	4.7	4.84	4.77	**	**	-	-	-	-	-	-	-	-
NT2RM4002076	5.05	4.42	4.84	7.24	1.99	2.02	1.65	2.18	1.6	**	**	-	-	-	-	-	-	-	-
NT2RM4002078	61.79	41.41	62.13	44.43	23.08	49.45	46.13	89.55	78.59										
NT2RM4002081	12.47	11.63	21.62	11.3	5.48	8.38	4.38	5.75	7.08	*		-	-	-	-	-	-	-	-
NT2RM4002082	28.78	23.31	28.71	31.25	27.4	16.48	17.58	15.02	15.49	**	*	-	-	-	-	-	-	-	-
NT2RM4002093	9.61	9.51	12.28	11.79	5.58	6.25	5.84	7.44	5.28	*		-	-	-	-	-	-	-	-
NT2RM4002109	6.91	6.93	10.41	8.11	5.84	4.18	3.3	3.14	2.63	*		-	-	-	-	-	-	-	-
NT2RM4002115	9.41	7.3	7.03	6.27	4.68	1.89	3.74	1.98	2.7	**	**	-	-	-	-	-	-	-	-
NT2RM4002118	37.45	58.41	50.21	20.19	28.02	17.58	13.85	5.39	8.55	*	**	-	-	-	-	-	-	-	-
NT2RM4002128	8.66	5.7	4.74	7.73	3.31	3.74	6.08	4.03	5.02										
NT2RM4002137	10.72	7.46	13.29	9.5	3.57	7.97	5.85	8.61	6.51										
NT2RM4002139	4.03	2.83	9.32	5.97	2.53	3.54	2.11	4.88	1.91										
NT2RM4002140	13.52	11.7	18.78	14.09	10.76	8.24	10.87	13.55	10.92										
NT2RM4002145	9.85	17.52	15.65	7.49	7.31	6.12	9.29	8.25	6.8	*		-	-	-	-	-	-	-	-
NT2RM4002146	53.03	50.09	50.28	48.49	40.37	39.91	51.33	53.96	59.3	*		-	-	-	-	-	-	-	-
NT2RM4002161	42.85	50.59	52.58	30.41	24.15	28.58	27.02	24.86	21.9	**	**	-	-	-	-	-	-	-	-
NT2RM4002174	18.01	10.09	12.11	7.53	4.92	3.2	3.7	2.28	3.16	*	**	-	-	-	-	-	-	-	-
NT2RM4002178	12.55	13.35	13.21	15.19	8.38	11.5	9.87	9.09	6.99	**	**	-	-	-	-	-	-	-	-
NT2RM4002180	29.73	19.51	29	18.7	7.98	18.84	13.92	22.12	26.06										
NT2RM4002185	43.79	41.64	55.25	22.67	17.47	26.66	26.11	42.39	45.09	**	*	-	-	-	-	-	-	-	-
NT2RM4002183	7.67	8.99	10.29	5.98	5.23	2.74	5.92	2.8	1.53	*		-	-	-	-	-	-	-	-
NT2RM4002194	41.77	35.69	39.22	19.41	14.19	21.77	29.68	28.52	30.53	**	**	-	-	-	-	-	-	-	-
NT2RM4002198	85.18	103.48	83.23	24.68	23.75	38.65	24.56	34.17	38.68	**	*	-	-	-	-	-	-	-	-
NT2RM4002205	10.02	12.63	11.47	6.5	3.78	3.22	4.51	2.19	6.22	**	**	-	-	-	-	-	-	-	-
NT2RM4002213	21.88	23.22	24.09	13.45	7.91	12.62	20.89	17.99	21.68	**	**	-	-	-	-	-	-	-	-
NT2RM4002216	7.77	7.03	5.93	10.01	6.38	8.33	6.06	4.45	5.38										
NT2RM4002226	41.67	37.48	37.48	67.73	42.68	64.51	42.92	35.21	34.67										
NT2RM4002237	7.28	7.73	10.06	4.52	3.54	3.6	2.94	7.15	4.84	**		-	-	-	-	-	-	-	-

64 / 157

【0888】

【表417】

2567

2568

NT2R04002240	41.31	33.08	34.98	23.88	21.39	31.98	42.08	38.98	42.81
NT2R04002251	3.39	3.89	8.47	4.89	3.61	2.4	2.99	2.15	17.8
NT2R04002256	4.28	3.3	5.05	4.37	2.61	2.38	2.35	2.83	1.48
NT2R04002262	26.4	21.79	28.58	14.04	8.23	12.13	20.41	14.34	24.05
NT2R04002266	32	29.71	28.68	24.62	15.02	21.58	34.87	32.88	37.89
NT2R04002275	14.45	13.71	16.67	14.53	13.36	18.53	18.37	10.08	9
NT2R04002278	18.11	14.14	14.01	7.25	7.14	10.52	3.65	8.09	7.88
NT2R04002281	9.29	5.59	12.79	12.53	4.74	10.08	6.18	9.79	8.22
NT2R04002287	17.77	15.59	14.41	11.94	5.05	12.57	14.23	12.4	14.92
NT2R04002294	5.99	6.14	8.14	5.51	2.38	5.99	3.64	2.35	2.91
NT2R04002298	64.75	62.55	58.47	42.58	28.72	43.54	69.57	64.59	61.48
NT2R04002301	3.57	3.23	3.53	6.49	2.3	3.07	3.59	2.99	3.21
NT2R04002306	25.88	19.13	24.79	14.02	9.84	17.37	30.57	32.67	35.82
NT2R04002323	7.61	6.27	5.98	7.59	3.26	7.4	8.04	6.9	4.8
NT2R04002334	14.45	8.8	16.75	21.2	9.98	15.23	10.19	12.3	13.22
NT2R04002339	38.85	25.91	51.52	34.79	27.9	40.82	27.13	20.38	29.16
NT2R04002344	17.19	8.04	22.21	10.94	8.27	8.93	15.04	14.72	16.42
NT2R04002345	7.58	5.39	12.54	5.63	3.79	3.52	2.31	4.45	2.41
NT2R04002352	44.08	33.98	54.03	33.98	35.28	37.68	25.21	30.31	27.97
NT2R04002362	39.95	26.35	45.23	28.03	24.7	27.77	28.92	25.64	24.85
NT2R04002373	9.44	6.99	7.1	8.69	3.38	4.42	5.88	3.57	3.37
NT2R04002374	0.94	1.53	2.98	5.8	0.19	1.91	2.19	2.71	2.57
NT2R04002376	14.29	7.3	14.86	12.48	5.38	11.68	12.39	14.05	16.09
NT2R04002383	5.57	4.87	12.64	5.61	3.39	2.94	3.24	5.28	5.45
NT2R04002388	6.2	7.31	15.14	6.97	3.2	6.3	4.57	6.71	7.54
NT2R04002398	16.61	8.98	16.6	15.05	12.49	8.92	7.11	8.75	5.38
NT2R04002409	3.91	1.86	5.55	4.87	1.55	2.44	2.22	2.93	0.87
NT2R04002414	18.88	15.16	32.88	21.65	28.59	21.32	20.38	19.06	17.22
NT2R04002438	3.98	4.99	8.08	7.18	3.61	4.84	4.85	4.1	2.07
NT2R04002440	32.39	17.01	28.92	38.22	20.77	30.03	30.94	26.11	25.22
NT2R04002448	10.16	7.87	12.29	14.28	6.56	8.04	5.21	8.63	33.19
NT2R04002450	5.69	5.77	10.55	7.51	2.82	4.53	1.81	3.63	7.42
NT2R04002452	15.3	12.15	20.1	12.34	7.95	8.52	9.04	11.33	9.95
NT2R04002457	5.93	4.98	9.2	5.61	3.34	2.74	3.42	3.45	3.16
NT2R04002458	5.3	7.18	9	7.94	5.54	4.54	4.93	4.05	3.12
NT2R04002460	5.82	8.3	10.61	8.18	6.55	6.42	6.02	6.25	3.91
NT2R04002464	5.03	6.81	7.71	5.37	2.82	2.94	2.52	3.29	2.23
NT2R04002478	40.28	37.05	46.08	41.81	24.81	36.24	37.84	34.54	34.16
NT2R04002482	13.68	18.31	28.18	15.72	10.35	15.81	9.15	8.99	18.35
NT2R04002489	15.44	12.03	20.84	12.49	8.35	8.73	7.84	10.28	14.04
NT2R04002493	8.53	4.66	11.35	5.77	2.46	3.69	1.55	3.88	3.31
NT2R04002499	16.96	16.44	28.91	11.24	8.53	18.09	18.09	16.97	14.06
NT2R04002504	24.74	20.57	17.4	17.01	12.22	12.67	5.73	6.11	7.83
NT2R04002508	9.92	11.26	12.04	6.45	4.68	4.22	2.88	4.99	2.88
NT2R04002510	17.14	20.34	27.62	13.7	8.64	11	8.3	5.68	5.29
NT2R04002527	5.27	5.32	5.24	8.73	2.73	4.8	7.63	6.1	4.65
NT2R04002532	2.04	3.38	12.81	3.92	2.01	0.75	2.59	1.83	2.16
NT2R04002534	6.24	4.2	12.1	8.74	2.8	3.26	1.86	3.47	2.45
NT2R04002535	9.02	8.87	16.41	7.91	3.99	2.92	0.97	4.02	2.94
NT2R04002554	6.47	7.38	8.92	4.87	2.87	2.20	2.04	1.83	1.15
NT2R04002558	10.72	10.21	14.55	8.7	6.67	5.1	5.89	7.82	7.9
NT2R04002565	7.12	9.02	9.28	6.37	4.46	3.35	5.55	4.05	6.48
NT2R04002567	8.35	8.05	12.48	6.62	4.6	3.2	2.87	5.5	6.18
NT2R04002571	3.38	3.03	4.55	5.3	2.26	1.53	2.86	1.83	1.84
NT2R04002572	3.42	1.92	7.29	5.02	1.66	0.9	0.78	0.74	1.09
NT2R04002577	17.78	12.61	20.74	8.52	7.02	10.13	3.18	8.47	5.97
NT2R04002583	26.66	37.71	48	24.28	17.87	13.18	6.08	13.68	11.34
NT2R04002584	24.65	25.28	41	25.45	15.38	14.04	7.11	11.26	5.72
NT2R04002592	13.18	11.32	14.62	7.42	4.85	4.9	3.83	4.05	2.43
NT2R04002594	17.08	15.66	20.52	11.9	10.98	5.12	3.63	4.19	2.45
NT2R04002604	8.24	6.68	8.88	5.84	3.74	2.95	2.65	0.99	1.52
NT2R04002614	14.57	8.99	9.81	12.08	5.4	4.41	5.49	5.87	6.71
NT2R04002616	31.41	23.47	38.09	26.19	21.94	4.64	13.84	17.64	27
NT2R04002623	6.22	6.78	13.92	7.63	3.69	4.81	3.87	5.8	5.63

65 / 157

2569

2570

NT2RM4002634	8.22	4.99	11.16	9.11	4.37	3.31	1.21	3.45	2.39			
NT2RM4002635	8.59	6.68	10.88	5.53	3.08	3.09	2.42	3.32	2.29			
NT2RP10000002	38.84	33.13	45.9	28.04	32.42	26.61	13.28	16.85	17.96	00	-	-
NT2RP10000003	16.16	12.37	15.85	10.42	5.99	5.72	9.46	7.81	8.28	0	00	-
NT2RP10000015	7.32	9.13	12.92	6.24	3.84	1.22	4.28	2.29	4.19	0	-	-
NT2RP10000016	7.99	5.64	4.74	10.16	3.74	4.75	7.93	7.02	6.47			
NT2RP10000034	93.68	89.94	88.11	64.65	53.66	88	72.54	94.61	102.97			
NT2RP10000035	18.17	13.62	22.35	15.17	8.84	11.35	7.31	9.51	10.39	0	-	-
NT2RP10000040	16.4	12.21	23.34	12.38	8.64	8.99	5.04	9.39	4.11	0	-	-
NT2RP10000042	3.77	4.47	8.68	6.24	2.47	2.4	1.5	1.51	0.43	0	-	-
NT2RP10000048	11.21	13.33	14.61	8.92	8.14	5.61	8.58	7.29	5.06	0	-	-
NT2RP10000050	5.73	12.28	7.67	5.34	2.29	5.33	2.26	2.09	1.81	0	-	-
NT2RP10000055	13.77	17.86	17.27	9.41	8.14	4.91	5.82	3.33	2.85	00	-	-
NT2RP10000059	2.7	2.41	1.36	4.8	0.95	2.39	2.76	1.39	2.38			
NT2RP10000063	17.16	17.65	11.27	8.7	3.73	4.62	5.36	8.5	8.3	0	-	-
NT2RP10000065	29.14	23.7	38.49	16.13	18.83	12.51	21.73	27.58	32.66	0	-	-
NT2RP10000072	39.99	32.45	48.63	26.8	23.08	25.9	35.59	35.1	31.68	0	-	-
NT2RP10000073	30.81	35.89	43.46	17.05	9.38	10.22	9.08	9.67	9.54	00	00	-
NT2RP10000075	9.27	9.25	12.57	6.24	5.48	5.5	2.69	7.69	4.73	0	-	-
NT2RP10000079	6.69	6.1	5.38	5.53	2.26	2.02	2.99	2.23	1.19	00	-	-
NT2RP10000080	8.49	8.44	8.35	8.63	2.81	2.65	4.1	3.05	2.07	0	00	-
NT2RP10000086	8.5	9.4	9.89	12.4	14.28	6.82	7.93	8.49	7.97	0	-	-
NT2RP10000087	60.2	68.97	62.22	84.59	58.98	88.52	51.31	53.27	70.97			
NT2RP10000089	42.88	33.03	33.63	24.16	25.57	29.27	22.12	38.13	28.4	0	-	-
NT2RP10000090	36.01	34.5	38.9	42.34	38.64	45.67	28.7	26.11	29.28	0	00	-
NT2RP10001000	842.61	1642.2	1396.7	386.69	827.1	384.76	1509.9	896.88	783.21	0	-	-
NT2RP10001001	35.43	98.15	94.63	31.12	33.53	50.26	28.26	23.64	17.24			
NT2RP10001011	7.76	7.95	10.33	5.26	2.18	1.28	2.31	1.62	2.67	0	00	-
NT2RP10001012	23.37	50.82	42.64	24.98	11.61	20.16	18.5	13.38	13.37	0	-	-
NT2RP10001024	25.62	22.22	19.79	7.84	2.34	7.4	4.62	4.07	4.19	00	00	-
NT2RP10001025	10.72	8.67	12.07	7.14	2.95	7.79	4.48	7.44	4.25	0	-	-
NT2RP10001029	53.88	53.28	57.25	42.69	30.82	50.19	47.48	64.98	53.17			
NT2RP10001030	19.47	14.75	28.76	11.14	15.19	12.69	8.3	10.06	14.98			
NT2RP10001054	3.83	3.62	5.78	5.04	1.58	2.69	2	2.85	2.74			
NT2RP10001063	9.61	17.58	10	11.58	5.58	6.03	6.39	4.65	5.07	0	-	-
NT2RP10001070	4.63	4.62	7.7	8.31	2.3	2.01	2.67	3.94	2.2			
NT2RP10001074	9.99	10.63	9.3	8.41	2.75	3.81	5.44	4.09	3.96	00	00	-
NT2RP10001081	14.28	14.61	16.73	10.84	15.44	12.79	10.24	5.66	5.53	00	-	-
NT2RP10001091	12.96	10.86	7.82	16.4	11.07	15.31	6.1	13.72	9.91			
NT2RP10002020	24.16	14.94	24.74	18.17	11.69	9.79	7.42	9.13	15.3			
NT2RP10002039	9.53	5.6	7.62	9.29	2.89	11.94	8.48	12.01	11.59			
NT2RP10002043	6.22	4.43	4.68	7.85	5.29	6.41	2.34	3.31	5.03			
NT2RP10002055	12.37	12.16	15.41	16.34	7.48	13.06	9.35	10.82	12.98			
NT2RP10002059	19.81	8.35	10.69	9.6	2.44	9.12	13.53	11.28	13.39			
NT2RP10002081	8.39	2.68	5.85	15.75	5.57	6.12	7.38	4.73	4.58			
NT2RP10002086	2.25	3.12	1.21	5.48	2.08	2.18	4.74	1.71	2.28			
NT2RP10002071	28.23	11.57	26.77	18.9	8.83	20.18	9.94	14.85	10.84			
NT2RP10002072	2.12	2.3	9.68	5.22	1.99	2.16	2.89	2.48	2.25			
NT2RP10002079	19.54	8.91	14.66	10.51	4.61	6.63	10.77	11.05	13.98			
NT2RP10002090	6.08	4.1	11.1	13.63	4.43	7.63	8.81	6.19	8.17			
NT2RP10002093	3.63	4.19	7.55	5.54	1.82	3.88	2.95	1.13	3.16			
NT2RP10003031	4.37	2.83	13.34	5.34	2.19	3.45	2.73	6.07	2.66			
NT2RP10003032	10.16	8.8	16.01	10.64	5.58	8.94	8.71	5.67	8.58			
NT2RP10003036	18.67	16.33	24.52	12.77	10.82	14.23	17.03	16.82	19.7			
NT2RP10003047	3.46	4.07	7.82	6.45	3	4.66	4.14	3.03	3.73			
NT2RP10003048	24.17	30.38	47.03	19.99	17.54	16.37	14.54	7.37	12.73	00	00	-
NT2RP10003049	3.22	4.33	6.7	6.49	4.18	3.15	4.9	4.23	3.34			
NT2RP10003053	180.66	121.66	192.83	151.27	93.19	147.88	118.94	185.86	159.53			
NT2RP10003056	8.01	8.29	12.26	8.92	4.2	7.42	6.21	5.29	12.39			
NT2RP10003057	38.01	31.52	38.61	32.16	21.31	27.89	24.36	32.21	30.56			
NT2RP10003058	9.65	6.55	14.28	10.15	5.03	7.58	4.14	4.11	2.37	0	-	-
NT2RP10003060	11.08	7.25	16.85	18.78	9.48	8.41	8.09	8.21	9.08			
NT2RP10003063	37.28	30.39	52.28	41.73	39.62	39.87	33.79	24.41	33.64			
NT2RP10003076	6.26	6.27	9.53	5.76	4.97	3.76	4.87	4.58	3.63			

66 / 157

2571

2572

WT2RP1000384	37.77	48.73	61.62	24.07	21.15	20.09	24.61	14.84	16.66	*	**	-
WT2RP1000407	20.7	11.19	17.51	22.94	11.29	15.13	8.7	15.9	29.09	*	-	-
WT2RP1000409	24.93	20.8	32.63	23.46	10.36	22.79	11.36	15.76	11.35	*	-	-
WT2RP1000413	24.89	20.12	32.27	23.23	14.81	20.34	6.71	17.88	15.72	*	-	-
WT2RP1000416	9.92	9.55	19.32	9.88	6.1	6.11	5.35	8.54	6.35	*	-	-
WT2RP1000418	5.47	5.48	8.68	6.04	3.64	3.96	4.07	4.76	2.23	*	-	-
WT2RP1000420	22.79	21.95	33.37	16.39	18.64	13.03	9.29	9.88	11.08	*	-	-
WT2RP1000434	13.91	15.33	17.37	9.5	9.94	9.87	16.13	15.05	16.89	**	-	-
WT2RP1000439	25.29	35.1	29.04	26.55	26.88	19.87	29.22	23.74	23.38	*	-	-
WT2RP1000443	20.2	15.54	27.59	12.04	8.66	7.66	5.79	6.84	10.23	*	-	-
WT2RP1000447	8.08	15.72	18.03	9.49	4.04	7.9	4.98	7.89	18.07	*	-	-
WT2RP1000448	3.96	4.08	14.01	5.34	1.34	3.27	1.83	3.09	2.48	*	-	-
WT2RP1000451	4.39	7.13	7.39	6.28	2.08	2.05	2.16	2.95	1.78	*	-	-
WT2RP1000456	7.7	7.09	11.38	7.08	4.08	2.49	4.41	2.72	2.94	*	-	-
WT2RP1000455	3.88	6.63	8.77	6.04	2.34	3	3.32	3.09	3.78	*	-	-
WT2RP1000458	10.59	11.91	14.42	9.41	9.31	7.51	8.52	6	7.58	*	-	-
WT2RP1000470	5.79	2.60	10.41	5.27	1.59	4.02	1.42	2.61	2.6	*	-	-
WT2RP1000477	82.32	107.32	97.78	75.25	82.97	57.64	20.6	83.67	85.18	*	-	-
WT2RP1000478	15.37	19.72	31.56	11.53	8.67	8.62	5.74	7.51	6.18	*	-	-
WT2RP1000481	26.31	30.78	29.03	24.24	25.58	11.39	8.04	10.78	12.11	**	-	-
WT2RP1000493	20.43	14.94	23.34	18.48	8.49	8.21	13.43	15.08	18.66	*	-	-
WT2RP1000513	46.32	35.58	52.07	60.82	43.2	48.5	28.73	37.16	31.64	*	-	-
WT2RP1000522	31.35	42.08	42.78	21.13	28.37	20.27	18.09	8.09	12.01	*	-	-
WT2RP1000533	3.26	4.01	2.18	4.33	1.86	1.15	1.34	1	0.91	*	-	-
WT2RP1000544	15.77	9.6	19.95	8.16	5.29	4.62	4.67	9.69	8.27	*	-	-
WT2RP1000547	6.82	3.97	11.11	7.34	2.91	3.12	1.8	4.62	2.98	*	-	-
WT2RP1000551	7.67	6.25	20.26	6.79	4.61	3.98	3.69	5.05	3.29	*	-	-
WT2RP1000567	7.6	7.79	12.24	11.47	5.44	5.88	2.72	3.24	2.13	*	-	-
WT2RP1000574	10.39	7.63	14.91	6.86	4.35	2.94	2.41	2.6	3.1	*	-	-
WT2RP1000577	6.72	6.64	7.37	6.89	2.31	2.08	2	3.14	1.89	**	-	-
WT2RP1000579	6.69	4.72	10.08	6.29	2.62	2.74	2.3	1.65	2.85	*	-	-
WT2RP1000581	11.66	12.4	16.41	11.21	8.47	7.93	11.76	7.04	4.88	*	-	-
WT2RP1000593	11.3	18.39	27.51	8.2	5.55	6.03	1.75	7.5	4.02	*	-	-
WT2RP1000604	24.51	26.11	37.01	27.05	13.43	25.97	13.69	26.68	23.37	*	-	-
WT2RP1000609	11.66	13.43	21.36	13.95	9.13	8.99	3.92	12.37	8.18	*	-	-
WT2RP1000613	17.95	20.62	29.59	16.28	14.93	12.95	12.7	12.66	9.93	*	-	-
WT2RP1000622	5.38	3.34	9.13	5.39	2.07	1.13	2.51	2.21	0.46	**	-	-
WT2RP1000627	4.69	3.23	6.07	4.67	2.08	1.44	1.45	1.18	1.68	*	-	-
WT2RP1000628	8.08	7.69	11.36	7.12	3.71	2.97	5.95	2.89	3.69	*	-	-
WT2RP1000630	7.73	9.3	8.35	7.9	4.49	4.92	4.54	3.4	2.81	**	-	-
WT2RP1000639	16.2	13.68	16.62	7.82	5.84	8.02	4.83	7.17	5.87	**	-	-
WT2RP1000640	7.36	9.59	17.48	9.53	6.74	8.49	5.28	6.04	7.68	*	-	-
WT2RP1000645	4.24	5.61	12.04	6.49	2.96	3.77	4.79	5.79	2.35	*	-	-
WT2RP1000659	23.32	18.8	20.24	8.66	11.89	4.68	4.8	5.33	5.98	**	-	-
WT2RP1000674	5.93	7.17	11.83	6.23	3.45	3.48	1.76	2.68	3.67	*	-	-
WT2RP1000677	14.6	24.32	25.71	10.51	12.87	7.3	5.18	10.1	7.28	*	-	-
WT2RP1000679	49.27	69.62	36.95	29.7	31.01	14.48	34.6	17.18	10.04	*	-	-
WT2RP1000686	8.01	8.65	7.8	7.2	2.8	2.42	6.02	2.46	2.38	*	-	-
WT2RP1000689	32.73	30.66	30.93	19.81	8.55	17.9	17.63	24.88	21.39	*	-	-
WT2RP1000695	17.35	14.65	15.87	10.45	3.47	6.36	2.82	6.77	7.19	*	-	-
WT2RP1000701	7.19	4.87	10.84	7.55	5.48	2.64	2.6	6.76	2.45	*	-	-
WT2RP1000702	7.2	10.25	10.38	6.65	2.25	1.64	3.18	3.24	2.79	*	-	-
WT2RP1000713	7.21	6.48	7.91	8.68	2.67	3.12	4.23	3.89	1.7	*	-	-
WT2RP1000721	16.01	8.63	33.56	6.6	3.3	4.77	2.96	3.62	2.32	*	-	-
WT2RP1000730	34.22	37.43	42.67	27.42	15.22	17.28	23.72	15.85	27.4	*	-	-
WT2RP1000733	9.95	10.41	8.91	8.92	3.88	6.5	7.94	2.92	6.71	*	-	-
WT2RP1000738	1.62	1.49	4.67	6.56	1.13	3.01	1.45	2.92	1.28	*	-	-
WT2RP1000739	19.87	17.5	27.32	13.78	6.59	15.22	15.07	28.31	27.46	*	-	-
WT2RP1000740	10.94	13.28	12.49	8.2	7.61	6.64	6.43	11.92	8.57	**	-	-
WT2RP1000746	3.62	5.92	7.73	5.84	4.22	2.64	2.78	2.98	2.29	*	-	-
WT2RP1000750	12.79	9.22	13.58	7.87	4.28	4.14	7.19	4.29	4.16	*	-	-
WT2RP1000751	10.42	10.46	12.4	8.67	5.65	4.98	4.2	5.02	6.07	*	-	-
WT2RP1000767	10.48	7.93	10.97	6.88	2.38	2.64	1.45	2.21	3.81	*	-	-
WT2RP1000769	4.03	4.96	3.53	5.29	3.09	2.68	3.04	5.45	2.56	*	-	-

67 / 157

【0891】

【表420】

2573

2574

NT2RP1000780	19.08	11.29	17.88	11.55	12.19	16.38	7.6	12.92	8.98				
NT2RP1000782	3.8	1.98	4.82	5.26	2.11	4.2	3.73	3.14	3.38				
NT2RP1000785	23.21	12.88	16.72	13.64	7.34	8.03	3.18	8.05	3.72	*	-		
NT2RP1000787	8.11	7.21	12.68	6.28	1.88	5.78	3.95	3.02	3.35	*	-		
NT2RP1000800	6.38	7.42	8.45	6.35	6.28	4.58	5.14	4.58	6.09	*	-		
NT2RP1000826	6.14	10.58	8.77	8.58	2.44	6.18	5.33	5.66	4.12				
NT2RP1000833	6.32	4.62	5.45	7.5	3.18	3.17	4.57	14.3	3				
NT2RP1000834	12.65	13.91	14.59	8.48	3.38	8.31	7.69	5.15	4.86	**	**	-	
NT2RP1000836	6.87	3.88	10.29	7.84	2.62	4.87	5.98	4.8	8.86				
NT2RP1000837	4.76	2.83	13.29	5.43	1.88	3.86	3.47	3.54	3.43				
NT2RP1000846	14.34	10.47	22.38	10.58	7.41	9.61	13.44	9.9	11.55				
NT2RP1000847	17.63	21.93	45.89	20.33	17.15	15.75	21.4	13.87	15.38				
NT2RP1000851	3.81	3.71	7.21	5	2.22	2.83	3.04	2.47	2.36				
NT2RP1000856	15.4	23.9	24.75	12.89	7.5	8.85	6.97	4.98	5.2	*	**	-	
NT2RP1000860	5.61	4.82	11.39	9.47	4.28	7.05	6.83	5.9	5.93				
NT2RP1000902	10.87	7.93	13.65	15.09	8.49	17.09	4.79	9.4	9.51				
NT2RP1000903	14.49	11.64	18.42	14.24	5.87	14.38	12.23	14.45	21.38				
NT2RP1000905	11.82	8.69	16.54	6.96	4.86	8.4	5.28	5.14	5.21	*	-		
NT2RP1000915	63.69	60.39	85.78	55.83	55.49	69.74	77.2	80.56	84.89				
NT2RP1000916	81.28	82.58	89.7	73.31	86.87	61.78	53.78	46.45	56.88	**	-		
NT2RP1000921	71.85	62.08	72.49	27.75	31.76	38.72	21.98	23.9	26.92	*	**	-	
NT2RP1000943	6.65	5.18	12.72	6.58	5.6	3.97	5.28	4.45	3.75				
NT2RP1000944	36.58	43.78	53.16	29.61	24.97	30.98	38.1	30.4	32.9	*	-		
NT2RP1000947	229.13	182.3	242.56	211.38	169.75	208.5	187.13	230.69	193.69				
NT2RP1000954	8.73	6.8	11.31	9.14	2.72	5.18	2.45	4.84	8.97				
NT2RP1000958	12.83	9.09	19.99	13.3	9.58	12.08	6.34	10.95	18.7				
NT2RP1000959	51.88	27.73	29.26	17.4	13.14	15.25	9.23	12.74	9.9	*	-		
NT2RP1000966	30.03	20.71	21.11	23.97	20.83	19.34	9.71	14.32	8.82	*	-		
NT2RP1000974	16.81	13.83	23.38	13.15	11.64	14.93	7.2	7.22	7.83	*	-		
NT2RP1000980	85.21	82.27	107.69	78.1	88.09	99.97	80	57.54	87.02				
NT2RP1000981	14.84	16.8	18.22	12.59	10.09	13.46	23.72	18.48	19.72	*	-		
NT2RP1000988	21.09	10.38	16.08	11.98	7.25	9.77	8.85	10.68	7.42				
NT2RP1001002	23.42	12.11	26.95	15.71	9.49	13.37	4.78	9.9	15.94				
NT2RP1001004	3.32	4.31	12.34	5.67	1.55	2.49	2.12	2.77	2.28				
NT2RP1001007	5.38	4.39	9.57	6.91	2.92	2.64	2.01	3.93	1.56				
NT2RP1001011	5.88	6.54	6.16	6.28	2.04	2.68	2.38	4.56	2.02	*	-		
NT2RP1001013	158.04	170.77	232.58	166.44	147.11	154.85	178.28	171.79	185.18				
NT2RP1001014	13.02	16.54	16.26	8.7	8	7.08	8.95	5.62	4.04	**	**	-	
NT2RP1001020	12.83	15.72	12.22	15.74	11.78	16.3	13.8	9.3	14.27				
NT2RP1001023	16.05	14.39	18.27	13.63	16.85	17.02	5.93	11.58	12.91				
NT2RP1001027	5.65	8.56	17.95	9.25	6.73	6.98	2.42	5.11	5.48				
NT2RP1001031	5.87	4.8	13.77	5.12	2.4	2.34	1.77	3.34	2.2				
NT2RP1001033	7.93	6.3	11.72	6.48	4.31	2.18	2.83	5.21	2.62	*	-		
NT2RP1001042	63.59	58.82	65.69	50.83	59.66	46.94	71.71	83.48	74.44	*	-		
NT2RP1001045	9.83	10.21	13.29	8.28	4.89	5.85	3.08	4.58	3.85	*	**	-	
NT2RP1001073	12.41	18.41	17.37	9.19	7.47	7.07	4.73	4.56	5.19	*	**	-	
NT2RP1001078	38.47	45.51	52.14	22.17	29.03	25.48	25.23	15.77	15.85	*	**	-	
NT2RP1001080	21.81	6.68	17.31	12.32	7.25	12.88	6.59	12.92	8.31				
NT2RP1001113	57.87	45.61	82.08	49.33	42.8	51.52	16.27	50.38	44.78				
NT2RP1001159	7.51	4.75	14.77	8.2	4.84	7.14	4.3	5.24	3.99				
NT2RP1001173	4.39	5.23	11.13	7.48	3.15	1.42	1.19	2.31	1.17				
NT2RP1001176	52.66	45.96	57.68	32.83	25.13	21.71	19.58	17.05	21.12	**	**	-	
NT2RP1001177	10.79	9.96	12.22	5.4	2.87	3.22	4.03	4.39	5.3	**	**	-	
NT2RP1001185	14.18	11.88	11.53	9.31	7.59	5.91	7.44	8.55	7.87	*	**	-	
NT2RP1001199	13.52	13.24	12.28	9.53	6.55	5.34	4.8	3.41	3.43	*	**	-	
NT2RP1001205	83.58	72.48	94.49	51.42	53.78	86.93	52.32	75.19	84.87				
NT2RP1001215	17.01	15.57	25.94	10.31	11.47	12.47	9.75	16.75	14.31				
NT2RP1001225	11.8	18.2	17.52	11.07	6.07	5.97	4.13	11.34	3.6	*	*	-	
NT2RP1001245	6.59	5.31	9.93	6.03	2.96	4.57	2.73	3.24	2.91	*	-		
NT2RP1001247	6.32	3.32	10.27	4.35	3.87	3.01	3.42	3.7	1.09				
NT2RP1001248	15.25	12.77	20.45	15.52	14.1	11.88	13.33	13.28	15.08				
NT2RP1001251	84.68	116.5	77.8	48.18	37.13	24.55	27.4	9.2	25.85	*	**	-	
NT2RP1001283	6.14	8.32	8.90	7.42	3.89	3.62	3.74	2.33	2.8	*	-		
NT2RP1001294	57.87	32.23	60.31	62.38	43.64	78.24	53.21	40.65	43.49				

68 / 157

【0892】

【表421】

2575

2576

WT2RP1001302	13.32	15.35	23.89	11.51	7.14	10.85	5.3	9.24	8.54	*	-
WT2RP1001310	78.58	77.83	107.95	40.93	38.87	35.03	13.95	16.52	15.88	**	-
WT2RP1001311	12.6	10.19	20.17	13.97	8.87	11.71	8.81	11.48	9.2	*	-
WT2RP1001313	47.33	35.38	51.8	18.55	17.75	13.75	4.5	6.14	3.62	**	-
WT2RP1001324	13.11	11.52	14.26	8.91	7.65	4.65	7	7.07	7.14	*	-
WT2RP1001349	8.69	8.34	8.01	6.09	3.98	4.38	3.75	1.87	1.49	**	-
WT2RP1001361	9.54	13.7	11.81	11.62	6.33	8.88	6.92	4.67	4.5	*	-
WT2RP1001379	13.14	11.82	15.33	13.1	6.47	9.99	7.99	14.61	8.34	*	-
WT2RP1001385	24.66	22.66	28.03	9.27	7.17	9.45	12.37	13.91	22.49	**	-
WT2RP1001395	5.65	15.28	10.88	10.51	4.01	1.7	3.74	3.38	2.71	*	-
WT2RP1001410	13.37	15.22	15.83	8.65	10.09	5.28	5.57	5.85	7.15	*	-
WT2RP1001424	28.73	25.67	34.05	32.08	24.11	29.55	22.85	15.99	24.49	*	-
WT2RP1001432	8.64	7.96	12.22	8.88	4.28	6.46	5.04	7.62	6.96	*	-
WT2RP1001449	19.34	21.85	20.51	10.44	8.38	4.06	8.88	5.88	6.23	**	-
WT2RP1001457	23.3	38.81	37.3	24.2	13.72	10.77	10.68	7.09	13.11	*	-
WT2RP1001459	13.65	11.78	16.02	12.39	4.68	8.62	6.06	9.05	6.34	*	-
WT2RP1001468	8.39	4.23	8.84	7.44	4.36	4.01	3.12	6.63	4.14	*	-
WT2RP1001475	11.89	17.85	12.68	11.05	10.38	6.52	5.05	12.24	6.54	*	-
WT2RP1001482	15.19	20.91	20.15	19.85	11.71	18.03	11.8	14.09	12.75	*	-
WT2RP1001494	3.95	5.83	10.18	5.63	2.66	3.18	2.73	3.52	2.73	*	-
WT2RP1001500	9.08	4.79	11.52	5.37	1.85	1.77	2.31	4.4	3.24	*	-
WT2RP1001517	18.13	18.45	18.1	8.4	4.89	3.69	7.15	5.48	8.44	**	-
WT2RP1001540	28.09	31.09	28.58	25.07	15.6	22.78	25.04	22.39	21.45	*	-
WT2RP1001543	29.19	16.33	24.14	16.6	10.89	13.4	7.07	16.07	9.64	*	-
WT2RP1001546	12.99	15.98	15.41	9.88	3.03	7.01	6.04	9.72	7.53	*	-
WT2RP1001550	8.32	14.7	14.31	7.95	15.82	5.57	6.4	9.76	8.41	*	-
WT2RP1001553	5.3	7.23	13.14	7.37	4.11	4.11	3.58	3.77	3.88	*	-
WT2RP1001555	54.78	52.8	55.48	31.87	21.44	34.31	43.17	41.32	32.44	**	-
WT2RP1001563	39.08	33.85	48.64	38.88	21.82	23.18	31.1	28.26	33.8	*	-
WT2RP1001589	3.88	1.89	1.05	5.34	1.73	1.64	1.86	1.23	2	*	-
WT2RP1001584	172.88	220.01	198.04	20.64	11.47	19.54	81.72	38.4	39.43	**	-
WT2RP1001599	4.12	3.82	5.18	7.25	3.83	4.87	3.81	3.84	3.39	*	-
WT2RP1001818	4.03	1.77	2.09	6.09	1.64	4	2.04	2.76	1.59	*	-
WT2RP1001854	14.62	9.64	12.33	11.97	7.7	10.53	5.54	7.93	4.12	*	-
WT2RP1001865	5.78	6.54	8.08	8.07	4.13	4.72	3.98	9.32	5.29	*	-
WT2RP1001879	8.69	7.12	7.88	9.21	3.08	4.08	4.04	6.22	5.1	*	-
WT2RP1001881	18.99	18.82	24.49	14.39	5.68	9.22	11.12	13.68	12.99	*	-
WT2RP1001894	8.52	3.33	5.98	7.17	2.61	3.23	4.87	4.47	3.87	*	-
WT2RP2000901	6.24	4.74	8.49	6.13	5.97	4.89	8.42	5.79	3.58	*	-
WT2RP2000905	8.03	5.67	13.4	11.28	7.14	5.76	7.62	8.34	6.39	*	-
WT2RP2000907	10.8	5.31	17.13	11.11	4.89	9.68	8.74	8.3	8.67	*	-
WT2RP2000908	5.65	3.68	9.46	7.12	2.84	3.68	3.74	5.6	5.37	*	-
WT2RP2000910	8.84	5.53	11.05	6.98	2.83	4.35	3.2	3.54	2.21	*	-
WT2RP2000911	9.38	13.48	17.28	10.79	8.33	9.28	5.31	3.23	5.5	*	-
WT2RP2000927	3.47	3.05	7.82	6.1	1.45	4.19	3.16	3	5.68	*	-
WT2RP2000928	4.74	6.32	6.74	8.05	2.13	4.67	3.21	1.86	4.03	*	-
WT2RP2000932	3.24	7.28	8.03	10.28	5.5	7.28	3.28	3.13	4.05	*	-
WT2RP2000940	15.29	8.48	28.48	11.5	6.56	9.38	4.88	6.35	7.25	*	-
WT2RP2000942	21.14	15.48	35.04	11.42	8.01	11.22	12.48	11.99	17.82	*	-
WT2RP2000945	5.83	3.94	13.32	7.43	3.18	3.46	3.52	5.2	3.95	*	-
WT2RP2000951	13.61	9.89	23.1	13.04	8.32	9.72	10.33	13.1	11.12	*	-
WT2RP2000954	16.51	12.89	19.57	11.83	9.96	11.02	9.84	7.65	7.91	*	-
WT2RP2000956	8.15	8.7	14.22	6.37	4.1	4.09	7.75	7.98	8.74	*	-
WT2RP2000957	7.7	5.49	9.31	7.63	5.41	5.64	12.9	3.34	4	*	-
WT2RP2000957	7.84	15.12	15.64	18.12	18.74	10.15	9.45	5.64	9.95	*	-
WT2RP2000970	522.83	440.99	557.3	484.06	449.31	610.82	418.05	682.99	650.83	*	-
WT2RP2000976	5.99	4.88	10.04	7.01	1.06	1.43	1.54	3.81	5.87	*	-
WT2RP2000977	14.65	10.4	23.49	9.32	7.18	6.28	8.55	8.84	8.57	*	-
WT2RP2000979	8.19	8.49	12.44	8.77	5.26	7.58	6.72	6.46	6.39	*	-
WT2RP2000983	17.05	20.32	26.56	22.32	20.65	23.39	19.38	14.35	14.55	*	-
WT2RP2000991	33.24	18.21	39.37	24.16	20.9	17.89	11.58	9.95	7.77	*	-
WT2RP2000992	0.22	2.2	2.32	4.37	1.08	1	2.25	1.85	1.49	*	-
WT2RP2000997	10.19	11.77	12.54	11.39	6.78	8.17	7.35	3.3	4.01	**	-
WT2RP2000998	14.34	13.24	18.96	13.26	5.07	9.1	5.48	9.38	18.63	*	-

69 / 157

2577

2578

NT2RP2000108	10.87	8.33	17.31	10.58	6.71	6.74	4.11	5.62	13.35				
NT2RP2000114	14.61	10.83	24.65	15.49	7.8	10.68	10.74	13.91	10.18				
NT2RP2000116	12.92	13.17	18.83	12.07	8.1	10.87	4.57	5.57	3.24	**	**	-	-
NT2RP2000119	50.32	47.2	56.58	31.02	20.71	31.82	10.15	20.31	20.57	**	**	-	-
NT2RP2000120	7.9	9.18	10.99	6.82	6.21	6.5	4.81	5.84	4.89	*	*	-	-
NT2RP2000126	27.18	24.32	32.55	18.42	11.34	17.52	24.37	18.13	19.51	*	*	-	-
NT2RP2000133	6.31	5.61	8.18	7.63	3.91	7.78	7.66	10.7	6.35				
NT2RP2000147	12.98	10.72	23.8	11.64	4.41	11.41	3.82	6.37	10.62				
NT2RP2000153	4.49	6.28	13.18	7.15	3.25	3.2	2.86	3.8	4.82				
NT2RP2000166	7.61	8.37	12.68	6.43	4.23	4.03	2.05	6.21	1.59				
NT2RP2000187	64.87	65.88	69.1	57.98	47.99	49.02	51.12	61.28	54.87				
NT2RP2000181	8.68	7.44	8.1	8.5	4.93	4.27	4.28	4.21	2.7	**		-	-
NT2RP2000186	8.64	8.81	11.55	8.08	4.98	11.13	4.05	3.85	9.74				
NT2RP2000173	5.87	12.11	13.07	7.31	10.58	7.74	13.29	16.36	4.1				
NT2RP2000175	8.28	6.9	5.57	9.33	4.37	3.63	6.8	5.65	4.1				
NT2RP2000178	7.4	9.25	18.07	7.99	4.34	5.36	2.6	5.71	4.5				
NT2RP2000183	5.74	7.89	15.89	7.74	4.42	5.84	3.67	6.88	4.04				
NT2RP2000195	16.12	12.54	17.88	18.63	13.32	13.8	5.12	8.05	4.54	**		-	-
NT2RP2000204	22.18	23.48	35.46	16.84	14.7	7.3	7.38	8.83	4.75	*		-	-
NT2RP2000205	308.18	263.17	270.79	153.75	186.84	122.03	44.13	69.43	57.15	**	**	-	-
NT2RP2000206	5.24	2.8	6.75	5.38	2.18	0.7	2.89	2.08	2.38				
NT2RP2000224	24.97	24.03	27.21	17.41	19.54	13.73	10.21	10.44	6.58	*	**	-	-
NT2RP2000230	26.52	31.48	20.02	29.78	28.28	14.04	27.93	14.94	32.18				
NT2RP2000231	102.91	95.51	182.32	51.89	45.81	56.22	28.1	58.89	105.21	**		-	-
NT2RP2000232	18.39	14.23	28.59	15.4	9.71	12.45	6.13	8.73	7.72	*		-	-
NT2RP2000233	7.64	7.96	15.89	7.58	4.07	3.39	2.14	3	2.28	*		-	-
NT2RP2000239	68.01	139	172.9	85.94	84.53	44.49	55.89	58.76	25.78				
NT2RP2000240	5.47	3.94	10.61	4.8	1.25	2.17	1.78	2.32	0.43				
NT2RP2000244	6.07	5.52	8.36	6.23	4.28	2.64	2.65	2.47	2.84	**		-	-
NT2RP2000254	18.57	14.82	18.18	13.37	8.16	8.91	7.89	5.94	4.14	*	**	-	-
NT2RP2000257	38.73	47.22	36.2	40.72	25.44	29.27	34.28	29.33	25.58				
NT2RP2000258	13.08	11.66	21.23	12.78	8.27	8.08	5.57	13.01	9.17				
NT2RP2000261	8.27	7.35	17.74	7.68	2.93	4.17	3.42	5.31	3.84				
NT2RP2000270	3.96	6.69	13.81	17.82	8.28	3.85	3.35	9.95	4.71				
NT2RP2000274	91.38	80.94	71.7	69.79	55.47	43.9	66.54	63.83	61.61	*	*	-	-
NT2RP2000277	4.35	3.88	5.68	5.38	1.67	1.79	1.78	1.55	2.23	**		-	-
NT2RP2000279	7.44	6.18	9.49	8.39	2.22	2.73	3.81	2.78	3.21	*		-	-
NT2RP2000282	5.12	8.14	7.43	6.14	1.91	2.31	3.36	2.12	4.13	*		-	-
NT2RP2000288	10.29	7.89	9.75	9.32	5.88	5.55	9.02	7.12	9.38				
NT2RP2000289	21.4	17.79	32.55	15.37	7.32	16.77	22.06	23.42	27.49				
NT2RP2000297	4.51	6.32	10.63	6.94	2.23	3.64	3.53	4.19	2.55				
NT2RP2000298	8.06	8.79	13.64	5.87	3.8	2.92	2.96	3.04	1.11	*	*	-	-
NT2RP2000310	80.52	91.37	87.72	32.16	27.21	29.87	20.91	14.48	20.59	**	**	-	-
NT2RP2000327	9.08	9.43	13.33	5.88	3.81	4.9	2.21	2.04	2.13	*	**	-	-
NT2RP2000328	21.82	17.34	15.84	8.37	8.84	6.89	5.88	4.83	2.63	**	**	-	-
NT2RP2000329	17.76	16.87	20.08	10.58	7.07	6.18	10.73	6.52	6.14	**	**	-	-
NT2RP2000333	8.83	7.98	9.31	8.74	1.78	4.67	8.04	7.67	6.48				
NT2RP2000337	28.88	28.84	31.77	17.74	6.47	12.94	6.8	13.35	15.87	*	**	-	-
NT2RP2000346	6.79	9.29	9.75	14.24	7.42	4.48	6.89	17.82	10.4				
NT2RP2000357	14.11	13.18	22.49	5.83	2.27	3.95	3.58	4.39	2.68	*	*	-	-
NT2RP2000358	9.23	5.13	8.18	5.34	2.02	3.61	4.07	3.18	2.94	*		-	-
NT2RP2000386	5.06	4.61	8.09	5.12	1.19	2.67	2.27	2.31	1.55	**		-	-
NT2RP2000389	29.84	28.23	39.9	17.03	12.11	15.49	15.78	7.98	15.06	*	*	-	-
NT2RP2000376	28.15	35.88	51.75	24.31	15.42	18.34	35.37	27.75	36.03				
NT2RP2000394	11.62	8.66	8.45	10.5	5.28	8.79	9.08	9.09	12.63				
NT2RP2000395	2.03	7.18	10.2	4.43	3.16	1.53	2.05	5.24	2.77				
NT2RP2000412	13.13	15.98	18.35	14.6	8.73	8.07	8.73	18.17	12.59				
NT2RP2000414	19.9	19.23	18.62	8.95	6.33	5.68	6.29	8.15	8.52	**	**	-	-
NT2RP2000420	5.15	7.34	11.45	10.83	6.03	5.33	5.71	4.93	7.78				
NT2RP2000422	10.72	8.28	11.8	8.9	4.28	5.31	4.96	4.78	6.17	*		-	-
NT2RP2000426	7.55	6.99	6.8	4.75	1.02	1.72	2.91	2.28	1.94	*	**	-	-
NT2RP2000428	8.74	4.22	5.73	4.38	1.17	2.38	8.21	2.73	3.16				
NT2RP2000438	12.17	11.38	11.77	10.89	3.58	5.65	4.51	2.36	2.83	**		-	-
NT2RP2000447	12.75	6.84	10.08	6.38	4.33	7.41	3.21	7.55	5.61				

70 / 157

2579

2580

NT2RP2000448	25.71	13.78	32.08	17.1	19.22	17.28	5.88	10.24	7.42	*	-
NT2RP2000459	9.24	8.95	9.52	9.52	4.14	4.38	2.69	8.53	3.6	*	-
NT2RP2000479	6.63	5.55	7.74	7.89	3.38	5.62	5.8	5.59	6.42	*	-
NT2RP2000498	11.27	6.84	9.95	5.98	2.48	3.08	4.73	2.88	2.18	*	-
NT2RP2000503	14.72	13.18	15.3	32.52	21.1	19.93	18.56	11.89	14.4	*	-
NT2RP2000510	7.33	5.8	8.13	5.52	2.28	3.46	4.2	6.04	2.87	*	-
NT2RP2000514	4.57	2.69	2.25	7.44	1.98	3.28	5.71	4	3.87	*	-
NT2RP2000516	8.64	4.62	8.19	7.51	4.34	8.37	3.47	4.35	4.95	*	-
NT2RP2000523	4.11	2.38	11.01	7.29	1.39	3.22	3.15	2.27	3.38	*	-
NT2RP2000533	2.68	1.37	7.63	4.83	1.22	2.09	0.79	2.38	0.48	*	-
NT2RP2000540	6.16	5.88	8.08	5.48	4.38	3.8	3.01	4.98	1.52	*	-
NT2RP2000547	7.42	7.44	10.91	6.02	3.13	4.89	2.88	4.52	3.39	*	-
NT2RP2000567	49.31	60.55	82.94	43.48	53.17	74.9	83.43	53.58	63.67	*	-
NT2RP2000558	18.68	19.95	27.42	20.89	14.13	19.17	24.64	17.82	18.81	*	-
NT2RP2000564	0.43	0.11	2.04	5.89	0.93	1.24	2.05	1.59	0.63	*	-
NT2RP2000565	13.78	7.23	14.74	12.39	9.94	11.42	3.9	7.3	8.83	*	-
NT2RP2000583	10.87	6.32	13.24	7.55	3.8	6.68	4.41	6.83	9.88	*	-
NT2RP2000591	7.05	5.42	12.75	6.46	3.08	3.75	3.89	6.69	2.85	*	-
NT2RP2000599	45.38	33.04	53.33	45.68	38.74	39.04	49.48	50.96	42.82	*	-
NT2RP2000601	5.8	2.35	8.28	6.01	2.72	4.74	3.53	6.5	4.71	*	-
NT2RP2000603	10.6	10.97	14.75	12.67	11.91	10.27	11.35	11.32	10.67	*	-
NT2RP2000610	16.79	27.41	21.69	12.88	12.08	9.82	7.92	6.77	7.19	*	-
NT2RP2000614	22.5	27.77	21.5	15.84	12.38	13.2	11.48	7.34	10.15	*	-
NT2RP2000616	15.8	10.73	17.88	14.04	10.65	8.28	8.53	13.65	10.31	*	-
NT2RP2000617	7	4.89	10.44	5.79	2.49	3.27	2.33	3.91	6.88	*	-
NT2RP2000623	30.93	27.18	42.39	23.25	19.11	22.34	22.03	24.05	17.67	*	-
NT2RP2000634	5.41	5.13	8.98	7.02	5.17	5.22	4.91	5.31	5.98	*	-
NT2RP2000636	6.22	7.05	7.98	5.06	2.33	3.65	3.42	4.84	2.62	*	-
NT2RP2000638	5.4	5.87	8.77	7.49	5.36	5.46	5.01	4.21	4.67	*	-
NT2RP2000644	6.08	11.67	13.8	11.21	5.35	7.7	6.1	5.24	4.14	*	-
NT2RP2000649	69.57	64.76	72	79.38	45.65	58.22	93.91	57.37	68.85	*	-
NT2RP2000652	14.9	10.94	49.84	10.88	5.91	11.87	6.2	6.28	32.14	*	-
NT2RP2000656	12.57	7.15	13.54	8.29	4.4	6.3	1.69	9.03	14.78	*	-
NT2RP2000658	5.45	4.52	10.88	5.11	2.13	3.91	1.81	2.12	1.71	*	-
NT2RP2000663	4.8	4.28	7.43	5.58	2.25	6.97	4.47	2.44	1.11	*	-
NT2RP2000664	10.9	7.61	12.36	9.42	3.68	8.64	13.44	10.25	9.81	*	-
NT2RP2000686	33.38	33.11	48.17	39.58	39	37.32	34.58	25.83	30.2	*	-
NT2RP2000678	6.12	9.03	9.25	9.8	7.57	6.65	8.86	6.02	5.34	*	-
NT2RP2000694	16.87	11.51	13.29	13.35	8.47	12.93	22.9	14.8	18.71	*	-
NT2RP2000704	2.84	3.9	14.85	6.19	2.87	7.29	2.88	6.45	4.78	*	-
NT2RP2000710	8.84	8.7	19.22	7.25	3.74	6.4	5.24	8.81	6.9	*	-
NT2RP2000712	5.84	5.25	16.28	7.12	3.27	6.34	2.82	4.15	2.23	*	-
NT2RP2000715	23.49	31.67	27.44	16.78	13.7	10.05	7.2	11.02	9.68	*	-
NT2RP2000720	14.6	11.95	17.99	9.87	7.04	6.8	9.67	8.11	6.87	*	-
NT2RP2000731	130.48	114.16	171.08	139.58	136.44	144.08	91.22	91.1	91.43	*	-
NT2RP2000738	10.56	11	13.93	9.81	7.28	5.89	9.06	5.03	5.15	*	-
NT2RP2000748	12.42	9.17	9.8	13.86	6.05	5.01	6.86	4.28	5.35	*	-
NT2RP2000749	11.23	7.84	14.78	7.12	4.44	8.82	1.43	4.83	4.58	*	-
NT2RP2000758	18.71	12.4	22.27	12.06	7.43	9.25	3.34	5.27	4.98	*	-
NT2RP2000764	8.83	9.65	15.17	9.88	4.61	8.48	4.78	3.89	4.68	*	-
NT2RP2000786	39.21	45.67	54.49	39.05	28.43	20.35	18.87	26.46	19.71	*	-
NT2RP2000777	37.43	31.1	43.71	20.28	16.78	21.16	3.56	8.61	6.24	*	-
NT2RP2000786	10.79	9.8	16.45	9.08	5.82	3.61	8.49	11.1	8.19	*	-
NT2RP2000793	8.86	7.43	14.72	5.59	4.46	3.27	3.41	3.35	2.38	*	-
NT2RP2000795	13.41	11.17	11.19	12.62	8.7	5.88	4.89	4.55	5.82	*	-
NT2RP2000809	9.51	8.68	16.69	9.15	5.7	10.07	8.67	9.07	9.06	*	-
NT2RP2000812	10.78	5.64	19.78	9.49	6.33	7.02	9.58	10.65	9.48	*	-
NT2RP2000814	2.73	2.98	11.46	6.79	1.46	2.03	1.53	1.11	1.73	*	-
NT2RP2000816	5.26	5.27	12.58	5.83	4.32	3.54	2.81	2.6	1.81	*	-
NT2RP2000818	64.67	62.94	68.69	58.85	32.29	31.82	38.98	34.96	38.14	*	-
NT2RP2000819	10.64	9.84	11.3	8.11	4.61	3.58	5.06	2.23	3.74	*	-
NT2RP2000841	19.27	13.42	17.68	15.7	9.96	11	12.88	9.6	13.79	*	-
NT2RP2000842	7.9	5.25	9.24	8.2	4.88	2.43	6.17	3.9	4.72	*	-
NT2RP2000845	1.64	3.58	10.06	6.25	3.24	6.16	2.58	3.32	3.23	*	-

71 / 157

2581

NT2RP2000881	9.23	2.37	9.61	12.6	5.57	10.29	2.95	6.5	2.03				
NT2RP2000882	8.89	4.15	10.09	8.75	3.85	3.81	3.32	7.64	2.82				
NT2RP2000883	6.48	7.22	10.47	6.68	3.69	2.44	2.83	2.13	2.32				
NT2RP2000884	6.84	8.43	12.75	7.7	5.81	3.85	6.32	3.41	5.03				
NT2RP2000885	14.72	12.47	10.22	6.37	7.06	6.25	6.3	4.12	3.45				
NT2RP2000886	7.83	5.78	8.48	5.87	3.5	4.08	5.51	1.9	1.89				
NT2RP2000887	7.08	7.41	7.6	9.95	3.31	4.88	6.74	6.38	7.22				
NT2RP2000888	17.23	15.65	19.14	9.83	11.03	7.29	5.69	13.37	8.87				
NT2RP2000889	4.85	5.4	12.51	5.58	2.98	2.75	2.51	3.48	1.84				
NT2RP2000890	32.12	38.82	41.75	20.34	22.26	19.68	28.05	24.64	29.99				
NT2RP2000891	8.65	4.28	8.98	8.63	2.06	1.49	2.39	2.3	2.44				
NT2RP2000892	8.82	7.9	11.5	5.58	3.81	3.8	2.18	2.83	1.58				
NT2RP2000893	5.64	8.03	16.9	6.17	2.43	2.74	1.92	2.25	3.58				
NT2RP2000894	11.75	9.04	10.68	6.59	3.84	2.64	4.69	3.69	5.68				
NT2RP2000895	4.05	2.88	2.8	5.91	0.94	1.41	2.34	1.56	2.95				
NT2RP2000896	58.85	65.2	74.74	64.15	40.74	92.64	72.54	97.39	89.84				
NT2RP2000897	5.19	3.92	9.39	8.08	5.68	3.54	3.01	9.75	7.5				
NT2RP2000898	3.85	2.51	4.33	4.79	3.19	4.55	1.82	2.45	2.12				
NT2RP2000899	39.38	38.27	32.68	33.06	40.49	26.38	33.61	26.04	27.01				
NT2RP2000900	9.33	8.58	9.52	8.72	1.47	7.24	8.83	9.44	11.63				
NT2RP2001024	12.74	7.82	18.87	8.22	2.48	2.6	4.96	3.99	3.75				
NT2RP2001025	38.71	37.69	41.88	39.58	15.8	30.84	34.47	25.69	33.82				
NT2RP2001026	10.56	11.08	9.18	9.7	2.35	6.59	6.08	3.75	3.17				
NT2RP2001027	29.77	24.74	30.19	17.85	11.75	29.98	20.42	28.95	23.24				
NT2RP2001044	111.16	103.9	116.81	65.77	42.98	31.59	30.2	74.24	42.92				
NT2RP2001069	16.47	11.38	12.03	9.34	5.04	8.68	12.51	8.41	8.95				
NT2RP2001085	12.87	18.54	21.09	9.8	8.83	8.58	6.23	8.54	8.28				
NT2RP2001087	18.27	13.06	18.78	9.25	5.28	3.58	5.5	6.23	8.68				
NT2RP2001070	4.55	6.03	7.25	5.14	2.9	2.2	3.37	2.83	1.95				
NT2RP2001081	11.96	15.94	13.87	8.19	3.81	5.26	5.58	4.16	8.83				
NT2RP2001087	65.9	61.9	78.08	39.7	16.73	50.09	38.08	21.24	22.35				
NT2RP2001084	12.2	8.99	11.02	14.81	10.72	9.62	11.02	11.03	11.05				
NT2RP2001119	4.18	4.58	7.07	10.85	1.99	7.07	4.12	3.83	2.63				
NT2RP2001127	11.38	13.12	3.9	16.08	4.34	7.73	5.38	7.85	3.42				
NT2RP2001133	5.67	4.14	6.88	10.97	6.13	8.58	8.68	4.41	3.17				
NT2RP2001137	49.8	41.1	45.81	39.26	21.06	24.92	25.21	45.63	17.8				
NT2RP2001142	5.27	5.53	5.08	5.3	2.28	3.88	2.41	3.58	3.78				
NT2RP2001149	6.92	5.81	5.45	13.38	3.82	8.27	6.19	4.78	4.91				
NT2RP2001168	8.12	7.49	5.85	8.34	5.85	5.09	5.44	4.63	5.1				
NT2RP2001173	4.74	3.23	8.28	7.11	2.4	3.67	4.81	6.54	6.37				
NT2RP2001174	44.53	21.39	37.72	44.99	18.88	48.31	52.67	53.84	55.67				
NT2RP2001184	7.2	4.99	10.33	8.93	4.09	3.88	4.88	4.85	5.6				
NT2RP2001186	188.27	171.83	278.21	277.29	158.05	217.9	310.45	339.15	334.54				
NT2RP2001200	5.84	6.42	11.03	6.08	6.9	5.52	4.08	2.49	4.45				
NT2RP2001218	12.41	10.49	14.66	10.34	6.86	7.23	9.11	8.8	8.23				
NT2RP2001223	18.09	12.28	12.04	13.71	9	9.77	10.94	7.88	9.72				
NT2RP2001225	295.34	151.71	253.15	264.52	155.79	205.15	225.76	302.68	279.91				
NT2RP2001227	9.29	9.83	15.24	10.83	5.91	8.02	6.71	6.54	8.43				
NT2RP2001232	6.25	9.05	13.72	6.11	9.52	4.37	3.08	5.35	5.73				
NT2RP2001233	17.2	11.64	28.2	12.08	8.02	7.48	8.68	8.89	8.82				
NT2RP2001245	42.01	30.78	58.72	22.72	58.34	32.95	29.15	20.95	32.88				
NT2RP2001246	5.12	5.82	8.66	6.91	4.33	3.68	3.64	7.01	3.2				
NT2RP2001268	76.85	63.92	101.42	58.4	109.5	90.32	72.85	73.19	77.59				
NT2RP2001270	10.64	12.74	14.24	12.74	10.24	11.95	10.58	9.82	10.94				
NT2RP2001276	23.31	16.76	30.97	21.15	16.68	10.08	14.57	15.7	14.88				
NT2RP2001277	24.91	10.73	18.18	16.47	8.03	6.24	22.15	25.09	49.12				
NT2RP2001290	8.95	7.82	10.48	8.48	7.41	5.73	5.82	9.83	10.81				
NT2RP2001295	14.82	7.48	15.13	8.28	5.62	4.48	5.33	6.4	4.82				
NT2RP2001297	8.15	6.87	10.81	9.38	7.32	4.92	4.17	4.98	6				
NT2RP2001301	47.34	44.45	63.71	42.51	36.25	33.89	33.81	29.8	26.52				
NT2RP2001312	8.43	10.98	14.02	7.43	5.35	5.08	6.81	4.48	4.32				
NT2RP2001327	20.88	28.57	34.54	16.53	15.03	15.84	11.1	20.84	9.85				
NT2RP2001328	10.32	3.77	9.25	9.2	3.81	6.74	4.44	5.58	7.15				
NT2RP2001341	3499.8	2809.8	4099.1	1384.9	1644.2	888.17	1201.7	2570.4	2392				

72 / 157

2582

2583

2584

NT2RP2001347	539.84	514.01	628.74	565.74	422.96	418.2	495.32	685.14	429.88				
NT2RP2001366	38.71	33.77	43.04	28.42	28.24	19.52	18.19	26.65	21.17	*	*	-	-
NT2RP2001370	8.65	6.18	12.14	10.82	6.77	5.51	3.49	4.36	4.96				
NT2RP2001381	20.33	22.01	25.11	15	9.62	10.21	8.87	12.31	12.9	**	**	-	-
NT2RP2001384	23.09	21.55	19.18	14.87	11.09	13.2	20	18.4	14.87	**		-	-
NT2RP2001391	19.47	21.48	20.82	19.07	15.36	11.27	15.83	12.63	10.03	*		-	-
NT2RP2001392	6.22	2.38	9.23	6.23	2.48	4.35	2.68	3.41	3.72				
NT2RP2001394	9	9.62	18.06	7.33	4.84	3.55	4.94	5.64	7.88				
NT2RP2001397	5.62	4.95	15.76	7.71	3.52	1.79	2.61	4.45	3.74				
NT2RP2001400	33.05	35.87	41.2	15.06	15.87	8.49	5.11	9.85	5.55	**	**	-	-
NT2RP2001408	22.02	23.86	22.82	11.45	12.19	9.24	10.98	11.38	13.8	**	**	-	-
NT2RP2001420	13.99	18.37	19.98	12.59	10.04	8.9	9.18	10.79	13.16	*	*	-	-
NT2RP2001423	138.22	181.12	189.38	120.84	140.16	121.74	142.58	124.54	162.09				
NT2RP2001427	11.99	17.22	18.25	9.35	7.9	6.74	7.44	6.11	8.28	*	**	-	-
NT2RP2001428	11.05	7.81	14.33	11.13	3.4	8	8.33	6.84	8.32				
NT2RP2001438	28.98	21.88	34.92	22.78	25.35	19.1	15.23	27.9	27.44				
NT2RP2001440	21.88	12.74	27.62	14.71	8.47	17.13	15.85	24.16	14.88				
NT2RP2001445	9	10.8	19.43	10.91	10.54	6.62	8.68	8.73	7.21				
NT2RP2001449	28.07	34.09	40.82	25.02	19.29	16.85	14.58	19.57	13.99	*	*	-	-
NT2RP2001450	20.8	28.19	27.38	18.17	18.42	8.85	15.87	22.32	18.66				
NT2RP2001487	77.4	117.05	74.08	41.14	94.21	47.44	48.02	38.8	29.18				
NT2RP2001489	12.38	11.13	18.21	9.61	8.91	7.88	6.18	5.85	4.91	*		-	-
NT2RP2001489	16.58	12.17	28.4	14.68	6.35	9.1	6.53	11.37	10.31	*		-	-
NT2RP2001495	5.39	5.05	10.71	5.62	3.39	3.05	2.02	2.99	3.13				
NT2RP2001499	31.27	46.77	45.33	29.35	33.83	17.6	16.09	21.38	15.2	*		-	-
NT2RP2001506	7.58	7.79	10.57	9.25	7.24	4.11	3.34	2.98	2.75	**		-	-
NT2RP2001508	10.21	8.68	19.01	9.32	5.78	3.48	10.19	10.43	18.1				
NT2RP2001511	18.18	10.38	14.04	8.08	6.96	6.65	4.43	6.31	7.52	*	*	-	-
NT2RP2001514	8.2	7.15	8.01	6.7	3.51	2.31	2.13	2.46	3.84	**		-	-
NT2RP2001520	8.17	6.6	9.18	8.28	6.18	3.08	5.09	4.32	4.95	*		-	-
NT2RP2001526	15.83	17.52	16.82	11.76	3.98	10.59	5.88	10.57	11.84	*	*	-	-
NT2RP2001529	18.96	9.11	22.28	13.03	8.09	10.02	6.2	8.22	8.57				
NT2RP2001535	14.16	10.38	18.6	12.98	11.79	8.28	7.74	11.37	11.72				
NT2RP2001538	8.11	6.47	12.78	7.87	5.34	3.77	4	4.36	2.58	*		-	-
NT2RP2001547	9.99	10.67	12.82	9.74	7.58	3.49	4.05	3.74	4.33	**		-	-
NT2RP2001560	11.5	10.74	9.02	8.82	5.56	7.18	6.87	5.68	4.21	*	*	-	-
NT2RP2001562	17.39	15.7	17.62	10.84	8.02	7.21	8.92	9.2	10.24	**	**	-	-
NT2RP2001585	14.26	13.81	10.49	11.53	7.07	4.68	7.24	4.21	4.54	**		-	-
NT2RP2001589	25.02	15.42	32.18	22.03	11.09	26.81	24.28	25.39	21.94				
NT2RP2001576	129.89	178.91	114.02	47.3	48.86	66.77	33.87	55.78	72.18	*	*	-	-
NT2RP2001581	20.4	22.7	22.23	13.8	17.88	10.64	10.57	16.71	15	*	*	-	-
NT2RP2001597	17.48	9.3	13	11.25	8.35	11.5	4.54	4.57	3.83	*		-	-
NT2RP2001601	378.46	416.73	382.11	182.76	196.04	131.77	252.12	219.45	367.42	**		-	-
NT2RP2001613	6.29	5.75	7.25	4.54	4.25	3.06	3.72	3.99	3.98	*	**	-	-
NT2RP2001628	20.46	20.27	25.19	14.18	13.5	11.94	11.98	13.91	12.32	**	**	-	-
NT2RP2001634	4.72	7.04	6	7.51	3.83	2.83	4.15	1.66	3.09	*		-	-
NT2RP2001635	20.18	15.77	22.18	15.58	7.6	10.65	10.91	15.44	14.53	*	*	-	-
NT2RP2001660	27.82	47.45	35.88	27.8	15.28	22.85	18.57	20.69	20.08	*		-	-
NT2RP2001662	141.98	137.52	151.45	151.4	118.43	138.78	154.6	188.28	148.8				
NT2RP2001663	102.28	119.38	140.87	48.03	51.07	39.15	80.45	80.81	83.55	**	*	-	-
NT2RP2001672	6.83	9.61	11.14	8.86	4.73	5.02	7.16	6.7	5.29				
NT2RP2001675	7.51	11.75	9.85	8.41	3.17	2.8	4.39	4.7	4.28	*	*	-	-
NT2RP2001677	18.83	22.14	19.19	10.84	6.34	6.91	9.72	6.83	7.68	**	**	-	-
NT2RP2001678	10.08	8.87	8.82	7.97	6.41	4.67	9.69	5.34	9.32	*		-	-
NT2RP2001683	88.93	103.45	116.49	185.82	102.44	134.62	102.34	95.16	65.68				
NT2RP2001699	36.15	36.2	44.95	18.8	7.95	13.41	6.7	9.44	13.11	**	**	-	-
NT2RP2001707	6.77	3.9	7.75	5.48	4.72	5.02	3.35	4.21	3.45				
NT2RP2001720	8.16	6.41	9.83	7.22	8.68	5.49	4.59	6.11	5.89				
NT2RP2001721	33.41	34.55	34.5	25.33	18.3	17.83	22.85	38.18	33.9	**		-	-
NT2RP2001740	64.22	76.89	65.2	34.74	42.49	31.73	26.06	24.24	41.05	**	**	-	-
NT2RP2001740	6.3	15.07	7.65	6.81	1.99	2.37	4.7	3.94	2.45				
NT2RP2001755	8.58	8.68	5.69	6.1	3.13	5.08	4.74	3.64	4.01	*		-	-
NT2RP2001762	4.42	4.43	4.81	6.08	1.37	4.5	3.19	3.5	2.83	**		-	-
NT2RP2001768	7.43	5.23	6.6	7.88	2.48	4.84	3.03	6.68	2.45				

73 / 157

2585

2586

NT2RP2001769	74.25	64.15	75.05	48.19	51.48	48.88	18.14	20.26	21.95	**	**	-	-
NT2RP2001784	6.1	6.84	9.79	8.92	5.74	5.64	4.78	4.07	4.10			-	-
NT2RP2001805	18.64	23.08	23.59	18.75	15.28	15.88	12.77	15.01	19.71			-	-
NT2RP2001813	12.2	15.41	15.49	8.88	4.21	6.14	9.29	6.92	9.66	*	*	-	-
NT2RP2001817	50.4	58.09	53.81	41.79	32.65	39.63	54.02	42.75	58.58	*	*	-	-
NT2RP2001818	7.27	4.11	3.7	5.91	1.64	5.16	5.71	4.24	6.18			-	-
NT2RP2001837	24.91	17.75	31.09	23.35	29.21	19.49	21.43	23.69	22.4			-	-
NT2RP2001839	18.58	14.52	27.11	17.12	15.87	19.02	13.8	13.73	15.93			-	-
NT2RP2001881	19.37	18.8	26.93	11.83	7.45	7.51	9.29	9.04	10.22	*	*	-	-
NT2RP2001883	8.35	8.88	14.85	9.01	6.82	4.27	3.19	6.52	4.2			-	-
NT2RP2001876	62.07	48.1	60.43	41.68	38.63	33.22	53.55	54.23	52.55	*	*	-	-
NT2RP2001878	10.64	8.11	18.99	8.71	6.81	6.18	8.15	6.28	8.38			-	-
NT2RP2001881	11.51	12.01	17.12	11.39	8.77	9.38	10.8	14.93	9.65			-	-
NT2RP2001883	15.67	6.93	15.2	18.32	7.55	11.74	13.41	14.75	12.06			-	-
NT2RP2001884	1443.8	767.12	607.01	701.8	386.77	733.69	199.74	492.38	1029.8			-	-
NT2RP2001885	88.63	81.18	119.19	85.34	88.67	72.27	80.49	93.38	87.15			-	-
NT2RP2001888	13.46	6.9	14.73	6.96	8.72	3.84	5.67	4.43	5			-	-
NT2RP2001900	20.73	14.58	24.69	18.79	8.43	10.74	17.28	16.1	17.71			-	-
NT2RP2001902	18.05	14.47	16.38	12.77	7.63	7.85	11	14.28	12.43			-	-
NT2RP2001907	9.79	12.89	12.82	8.11	8.3	4.57	8.52	7.83	7.63	*	*	-	-
NT2RP2001916	17.49	19.44	20.82	11.45	7.41	6.04	7.94	7.59	6.62	**	**	-	-
NT2RP2001921	13.67	9.05	20.68	11.35	6.81	9.21	6.21	10.48	17.99	*	*	-	-
NT2RP2001926	34.47	22.9	35.27	22.68	18.7	11.83	10.15	26.13	26.9	*	*	-	-
NT2RP2001932	36.39	16.3	35.12	23.36	20.81	17.87	10.85	17.5	22.37			-	-
NT2RP2001936	11.42	9.59	15.71	9.32	7.07	4.58	8.12	12.42	62.46			-	-
NT2RP2001943	14.71	11.45	18.63	11.8	23.77	7.53	8.68	10.79	7.88			-	-
NT2RP2001946	20.64	20.42	25.64	8.64	12.56	9.33	5.24	6.37	7.54	**	**	-	-
NT2RP2001947	6.28	10.5	6.57	7.88	5.98	1.91	4.38	3.2	4.16	*	*	-	-
NT2RP2001956	21.87	10.17	24.79	16.81	12.12	14.12	16.82	10.29	13.33			-	-
NT2RP2001969	201.65	149.79	101.42	176.35	143.62	157.42	143.78	183.78	182.57			-	-
NT2RP2001974	4.33	7.95	17.7	7.78	4.49	4.49	2.81	5.46	10.35			-	-
NT2RP2001978	14.35	12.6	20.81	12.02	5.57	5.03	3.1	3.58	4.91	**	**	-	-
NT2RP2001985	13.77	15.22	14.24	9.55	5.93	6.64	4.83	6.51	5.47	**	**	-	-
NT2RP2001991	37.12	36.77	50.48	37.18	44.84	28.4	28.39	20.28	24.2	*	*	-	-
NT2RP2001997	13.99	12.07	14.82	11.27	12.08	8.47	13.49	12.5	12.19			-	-
NT2RP2002015	3.95	7.2	5.18	7.61	4.81	2.95	3.72	6.25	2.81			-	-
NT2RP2002017	14.92	13.42	18.23	7.3	6.04	8.42	3.4	6.85	8.43	**	*	-	-
NT2RP2002025	39.84	48.8	88.43	33.02	35.2	30.44	23.31	31.13	31.97			-	-
NT2RP2002030	28.92	28.65	40.93	26.27	23.9	23.74	29.55	38.7	30.22			-	-
NT2RP2002032	41.77	39.39	41.34	31.44	32.7	25.79	16.28	26.82	17.93	**	**	-	-
NT2RP2002038	7.84	6.18	11.1	7.44	5.25	3.93	2.93	4.43	2.72	*	*	-	-
NT2RP2002041	63.15	38.11	58.94	48.43	54.38	37.89	33.68	26.24	20.79	*	*	-	-
NT2RP2002048	19.79	18.39	22.62	13.61	15.41	9.7	20.12	17.67	17.54	*	*	-	-
NT2RP2002047	15.98	12.18	14.55	9.42	12.66	9.04	8.28	8.52	3.77	*	*	-	-
NT2RP2002050	6.47	7.96	8.94	7.07	2.69	4.33	1.22	3.6	2.62	**	**	-	-
NT2RP2002052	16.18	10.28	17.67	10.58	8.94	8.28	5.44	8.53	9.43			-	-
NT2RP2002056	10.7	8.12	13.82	9.12	8.72	4.7	3.71	8.14	3.23			-	-
NT2RP2002060	7.22	6.32	13.21	7.18	3.78	3.46	2.87	4.32	2.18			-	-
NT2RP2002063	27.7	19.33	28	16.08	12.42	6.01	14.89	17.08	15.58	*	*	-	-
NT2RP2002066	14.12	10.25	16.69	11.69	8.3	5.49	7.88	7.37	6.79	*	*	-	-
NT2RP2002070	47.82	69.53	26.7	25.88	60.9	22.06	11.62	12.95	13.2	*	*	-	-
NT2RP2002076	50.79	54.44	68.91	34.7	35.7	26.78	21.75	13.08	12.15	**	**	-	-
NT2RP2002078	3.82	6.23	12.24	6.03	3.97	4.06	0.81	5.31	3.09			-	-
NT2RP2002079	13.88	8.15	17.98	9.69	6.98	8.2	5.55	8.29	8.41			-	-
NT2RP2002099	11.59	12.19	18.2	8.81	6.98	5.14	5.67	8.29	5.75	*	*	-	-
NT2RP2002105	15.09	12.54	15.38	10.34	7.26	6.01	5.6	9.12	6.68	*	**	-	-
NT2RP2002115	15.39	14.23	22.91	11.04	12.24	9.95	11	13.28	11.08			-	-
NT2RP2002124	59.88	68.27	77.47	56.69	47.35	38.88	55.21	47.08	46.87			-	-
NT2RP2002137	8.42	8.75	8.28	6.56	6.55	3.48	2.47	3.09	3.68	*	**	-	-
NT2RP2002139	93.62	131.25	91.98	52.77	52.45	52.95	81.88	52.45	48.22	*	*	-	-
NT2RP2002154	8.04	5.02	12.4	5.24	1.47	5.38	2.02	3.26	3.03			-	-
NT2RP2002165	16.04	17.28	22.84	17	10.94	11.7	12.12	14.76	13.07			-	-
NT2RP2002172	26.05	32.14	34.35	17.74	14.37	15.18	12.78	22.58	16.28	**	*	-	-
NT2RP2002185	13.73	14.4	21.2	14.18	14.86	10.72	12.32	12.52	11.2			-	-

74 / 157

2587

2588

NT2RP2002184	222.52	243.76	128.64	151.89	124.73	74.73	64.57	82.05	88.67	0	-
NT2RP2002192	68.59	140.59	123.86	61.74	74.83	59.88	111.82	74.79	70.77	0	-
NT2RP2002193	9.92	9.38	5.54	5.64	8.2	3.35	4.38	2.62	2.3	0	-
NT2RP2002208	24.07	21.97	28.87	14.79	10.85	11.7	14.39	13.9	16.1	00	-
NT2RP2002219	67.68	67.68	64.54	42.26	45.01	39.83	12.23	52.48	39.63	0	-
NT2RP2002231	7.82	8.91	9.85	5.33	3.41	3.41	3.43	5.73	5.85	00	-
NT2RP2002232	82.24	77.83	81.5	78.38	94.62	65.26	53.26	57.27	68.31	00	-
NT2RP2002235	28.18	33.42	30.04	28.58	26.18	22.07	20.58	21.22	22.01	00	-
NT2RP2002238	48.02	63.63	67.29	21.58	16.63	26.15	34.54	22.68	19.45	00	-
NT2RP2002252	7.05	9.07	9.23	8.29	3.79	3.9	4.57	3.45	2.91	00	-
NT2RP2002258	27.97	26.88	27.09	16.84	11.06	11.47	26.86	22.34	28.47	00	-
NT2RP2002267	7.18	9.4	7.62	6.89	3.78	4.59	5.58	4.76	6.16	0	-
NT2RP2002259	18.53	13.43	11.28	24.93	14.8	14.24	10.77	18.34	15.94	0	-
NT2RP2002264	28.23	33.28	22.8	12.91	13.58	12.04	9.7	17.76	13.48	00	-
NT2RP2002267	32.17	30.65	32.28	23.04	23.01	26.57	18.83	21.9	22.12	00	-
NT2RP2002270	24.56	42.02	39.93	12.58	23.51	8.23	20.01	17.31	11.5	0	-
NT2RP2002281	77.89	98.13	97.81	150.54	78.9	98.99	77.38	60.86	65.81	0	-
NT2RP2002288	21.39	25.5	23.86	18.7	13.7	14.18	19.97	22.13	17.3	0	-
NT2RP2002292	64.8	51.47	64.64	33.71	24.08	25.61	33.57	27.11	21.67	00	-
NT2RP2002299	38.27	50.42	61.77	19.48	10.3	12.83	10.02	9.38	6.36	00	-
NT2RP2002304	11.94	8.17	8.22	10.44	3.85	9.51	5.62	11.18	9.83	00	-
NT2RP2002312	27.46	21.25	33.88	24.99	23.29	27.25	22.52	32.81	25.31	0	-
NT2RP2002316	51.29	40.88	55.81	28.92	27.74	16.77	8.08	18.25	13.89	0	-
NT2RP2002315	14.79	13.26	29.38	13.29	7.49	7.83	6.29	6.54	6.53	0	-
NT2RP2002313	37.71	73.03	60.93	26.88	22.98	22.97	22.31	24.03	28.79	0	-
NT2RP2002371	18.65	14.9	20.81	15.68	7.83	11.84	19.13	18.45	18.7	0	-
NT2RP2002372	7.7	6.37	4.6	6.91	3.79	4.83	4.84	5.13	5.14	00	-
NT2RP2002381	9.55	13.3	13.28	6.59	3.77	3.7	6.54	4.62	6.35	00	-
NT2RP2002385	0.45	2.78	2.19	4.9	1.81	3.5	0.24	0.85	1.85	00	-
NT2RP2002394	32.72	28.71	26.62	28.1	25	26.16	15.67	15.8	20.33	00	-
NT2RP2002408	6.97	5.82	5.11	6.31	2.85	4.31	2.1	5.38	5.24	00	-
NT2RP2002409	7.43	7.51	7.75	10.28	4.99	7.17	4.61	5.21	4.74	00	-
NT2RP2002424	13.74	18.21	4.21	14.47	4.22	10.67	7.29	17.72	5.01	0	-
NT2RP2002426	31.08	22.01	25.45	29.33	10.8	28.89	38.27	34.35	51.45	0	-
NT2RP2002428	5.25	5.46	6.68	6.82	3.95	4.28	5.64	2.98	5.77	0	-
NT2RP2002437	5.68	8.15	5.2	8.99	3.78	4.37	5.91	5.88	5.8	0	-
NT2RP2002439	23.83	11.85	28.83	30	21.32	18.74	18.89	27.87	21.85	0	-
NT2RP2002442	4.21	2.48	10.08	7.13	2.17	3.83	3.89	4.18	3.7	0	-
NT2RP2002487	5.91	3.88	9.43	6.08	1.82	4.35	4.32	3.38	4.15	0	-
NT2RP2002484	14.04	9.04	19.39	8.28	2.91	3.82	6.2	7.3	8.31	0	-
NT2RP2002478	18.47	16.05	30.87	28.73	13.63	23.64	27.01	26.72	27.82	0	-
NT2RP2002479	4.38	4.25	7.88	9.08	3.28	4.87	3.38	7.1	3.51	0	-
NT2RP2002487	8.46	11.73	16.26	11.78	4.98	7.15	5.71	6.33	8.37	0	-
NT2RP2002488	12.42	14.56	14.64	14.3	15.74	13.48	11.49	11.5	14.04	0	-
NT2RP2002503	38.54	28.71	38.28	38.97	28.13	31.61	31.04	37.78	40.27	0	-
NT2RP2002504	10.68	11.45	20.66	9.47	10.7	9.29	7	7.64	9.65	0	-
NT2RP2002510	35.82	27.78	43.7	26.78	25.92	21.7	28.92	29.43	23.5	0	-
NT2RP2002520	21.45	19.39	27.79	18.61	13.24	8.38	14.05	12.04	8.59	0	-
NT2RP2002527	36.1	23.32	35.4	20.78	25.32	20.27	27.42	26.23	15.83	0	-
NT2RP2002513	13.17	14.75	23.11	11.93	9.58	7.87	8.62	7.98	9	0	-
NT2RP2002537	13.08	12.39	14.79	10.65	13.69	8	8.5	6.59	7.09	00	-
NT2RP2002542	68.44	60.08	72.33	74.8	48.95	45.75	49.02	38.47	37.92	00	-
NT2RP2002546	14.88	18.98	17.6	15.64	9.68	8.83	6.05	11.83	13.87	0	-
NT2RP2002549	14.98	11.64	18.89	10.43	8.78	6.03	5.02	8.52	8.5	0	-
NT2RP2002564	39.56	26.21	34.95	20.11	17.81	17.77	18.21	24.98	19.23	0	-
NT2RP2002591	42.59	28.9	37.43	32.97	20.53	25.24	26	30.71	25.2	0	-
NT2RP2002595	5.68	8	12.85	9.34	7.39	5.51	5.35	4.75	4.31	0	-
NT2RP2002602	54.53	65.87	75.68	69.84	57.23	46.71	59.72	54.59	54.85	0	-
NT2RP2002606	12.62	13.19	14.5	13.95	17.61	8.8	13.88	12.61	14.58	0	-
NT2RP2002609	11.6	17.33	13.31	20.12	19.43	14.55	10.05	12.64	13.08	0	-
NT2RP2002618	14	10.22	18.04	13.28	7.51	9.59	7.58	12.06	11.33	0	-
NT2RP2002621	11.65	9.77	15.54	8.04	5.31	5.01	3.35	5.97	7.64	0	-
NT2RP2002643	11.04	8.19	16.62	9.29	6.05	6.59	3.63	6.63	4.9	0	-
NT2RP2002672	6.88	6.19	9.33	7.99	5.69	4.36	3.64	6.54	3.29	0	-

75 / 157

2589

2590

NT2RP2002673	103.76	111.24	114.16	70.07	45.09	85.59	92.44	85.88	91.14	*	**	-	-
NT2RP2002674	28.14	27.69	34.14	21.08	20.78	18.49	24.37	22.8	22.81	*	*	-	-
NT2RP2002686	20.76	27.04	31.72	23.28	22.04	15.33	8.73	8.68	8.13	*	**	-	-
NT2RP2002688	13.92	12.24	10	14.56	10.68	8.28	14.43	8.67	10.91	*	*	-	-
NT2RP2002696	10.25	14.55	23.17	14.76	11.8	15.12	11.5	15.77	21.62	*	*	-	-
NT2RP2002701	4.76	5.4	12.91	8.52	3.02	6.45	3.43	3.42	5.6	*	*	-	-
NT2RP2002706	14.53	15.48	22.22	9.32	11.09	8.76	5.94	11.12	9.18	*	*	-	-
NT2RP2002710	30.81	28.84	37.75	21.5	19.35	13.67	11.45	15.23	13.68	*	**	-	-
NT2RP2002721	7.29	8.84	7.35	5.81	4.84	2.99	3.61	3.98	1.82	*	**	-	-
NT2RP2002727	6.54	9.38	10.59	7.9	7.44	4.24	4.08	5.05	9.6	*	*	-	-
NT2RP2002734	48.13	41.34	40.09	29.15	43.09	31.56	39.87	34.2	33.11	*	*	-	-
NT2RP2002736	8.71	9.47	8.91	9.58	7.4	9.48	7.84	7.21	5.83	*	*	-	-
NT2RP2002740	19.4	13.22	23.87	11.87	7.31	11.43	6.19	11.24	13.39	*	*	-	-
NT2RP2002741	23.72	17.52	33.51	16.7	12.26	14.24	12.67	22.66	22.7	*	*	-	-
NT2RP2002750	63.6	50.03	64.42	39.1	24.44	35.13	38.13	36.29	30.68	*	**	-	-
NT2RP2002752	10.52	9.47	15.12	8.58	15.15	3.88	2.6	5.03	3.54	*	*	-	-
NT2RP2002753	12.15	11.11	17.25	9.74	8.82	4.65	6.04	8.44	4.68	*	*	-	-
NT2RP2002760	73.48	81.4	101.38	101.02	83.73	82.2	53.54	64.02	59.41	*	*	-	-
NT2RP2002769	82.25	85.07	85.68	43.44	87.23	54.77	13.8	14.78	19.89	*	**	-	-
NT2RP2002778	10.58	7.74	8.45	8.8	4.88	4.14	4.05	4.2	5.19	*	**	-	-
NT2RP2002791	21.86	17.89	24.67	15.97	10.79	19.44	8.98	15.98	17.2	*	*	-	-
NT2RP2002800	57.56	44.66	67.26	27.41	29.24	33	38.77	66.85	81.38	*	*	-	-
NT2RP2002805	10.55	19.25	27.64	13.98	23.59	9.88	7.4	10.07	7.75	*	*	-	-
NT2RP2002811	13.01	13.43	18.61	12.19	7.67	8.64	7.16	10.12	7.7	*	*	-	-
NT2RP2002824	16.53	12.33	20.93	15.22	10.57	11.14	9.08	9.37	7.84	*	*	-	-
NT2RP2002839	7.81	6.4	5.94	5.9	4.06	3.3	9.77	3.03	3.13	*	*	-	-
NT2RP2002845	12.27	9.25	11.68	10.47	8.08	5	6.84	6.96	4.08	*	*	-	-
NT2RP2002857	13.78	18.16	14.57	14.66	30.41	11.09	9.34	6.45	5.36	*	**	-	-
NT2RP2002882	39.8	48.81	41.43	30.51	20.27	24.78	25.07	37.41	27.22	*	*	-	-
NT2RP2002883	68.2	85.99	77.82	51.13	34.35	43.94	48.02	65.53	68.81	*	*	-	-
NT2RP2002885	6.6	7.46	15.43	8.27	7.08	4.47	4.01	3.85	3.27	*	*	-	-
NT2RP2002891	7.27	7.43	13.2	9.04	9.22	4.71	4.88	7.24	5.8	*	*	-	-
NT2RP2002907	27.84	27.28	29.25	27.92	20.22	19.34	24.71	20.15	30.82	*	**	-	-
NT2RP2002925	8.14	12.06	8.92	8.22	5.24	3.39	2.57	2.62	2.32	*	**	-	-
NT2RP2002927	11.04	11.63	9.75	8.5	11.71	5	4.88	3.47	4.87	*	**	-	-
NT2RP2002928	14.93	19.65	15.07	18.82	17.04	17.98	11.05	6.88	7.13	*	*	-	-
NT2RP2002929	34.38	37.86	32.48	20.14	17.6	22.45	18.17	27.25	31.67	*	**	-	-
NT2RP2002934	25.44	20.87	30.89	18.19	13.85	16.46	17.37	22.76	21.39	*	*	-	-
NT2RP2002935	16.49	24.31	21.25	14.42	15.1	11.79	7.22	8.55	7.99	*	**	-	-
NT2RP2002942	60.99	60.19	88.43	63.29	72.60	51.11	69.34	65.39	67.68	*	*	-	-
NT2RP2002954	14.37	21.3	26.97	15.75	10.45	11.22	8.24	8.15	9.35	*	*	-	-
NT2RP2002959	11.9	10.85	14.73	7.3	8.01	4.57	5.17	3.69	3.58	*	**	-	-
NT2RP2002974	27.97	23.95	30.02	18.63	11.35	13.46	21.06	14.1	16.2	*	*	-	-
NT2RP2002975	33.37	27.08	30.74	41.03	19.52	40.87	48.18	35.03	48.85	*	*	-	-
NT2RP2002979	34.44	28.83	35.03	24.79	13.64	25.53	28.21	33.42	38.67	*	*	-	-
NT2RP2002980	58.79	52.53	62.74	87.66	81.62	49.88	38.91	45.4	40.62	*	*	-	-
NT2RP2002986	8.3	8.37	8.65	8.06	9.31	3.85	4.83	5.08	6.03	*	**	-	-
NT2RP2002987	103.6	100.12	88.59	73.68	56.64	94.83	89.25	107.73	109.15	*	*	-	-
NT2RP2002988	13.65	24.45	23.43	8.39	5.18	5.67	8.67	7.13	8.64	*	*	-	-
NT2RP2002989	30.07	31.12	32.88	20.82	20.33	19.14	25.73	28.88	29.98	*	**	-	-
NT2RP2003000	23.19	27.52	36.47	14.21	13.97	11.67	18.09	13.68	17.88	*	*	-	-
NT2RP2003008	20.38	21.66	17.83	30.65	18.58	30.92	20.83	11.64	23.71	*	*	-	-
NT2RP2003020	8.4	9.59	8.76	9.59	3.53	4.92	4.04	6.43	8.19	*	*	-	-
NT2RP2003032	16.62	21.12	22.97	11.47	10.11	7.97	7.7	9.72	7.64	*	**	-	-
NT2RP2003034	9.72	12.06	10.53	7.37	15.19	4.72	7.75	4.61	8.9	*	*	-	-
NT2RP2003042	7.27	6.68	12.91	7.34	5.07	3.54	6.69	5.74	4.38	*	*	-	-
NT2RP2003050	41.14	39.77	34.47	22.7	14.88	22.18	30.79	33.73	32.33	*	**	-	-
NT2RP2003060	13.62	10.73	6.38	6.73	19.61	3.28	5.01	3.28	4.49	*	*	-	-
NT2RP2003073	11.68	8.02	5.9	7.53	3.76	4.73	2.81	3.98	5.14	*	*	-	-
NT2RP2003098	15.8	18.78	12.88	19.38	10.86	18.98	13.15	11.19	10.41	*	*	-	-
NT2RP2003108	16.99	10.19	16.94	12.41	6.81	13.61	8.41	10.09	7.85	*	*	-	-
NT2RP2003116	140.62	119.44	125.07	149.82	108.25	164.27	145.7	152.43	149.71	*	*	-	-
NT2RP2003117	8.58	4.12	8.6	7.48	2.86	5.97	6.82	5.37	6.56	*	*	-	-
NT2RP2003121	15.37	11.98	21.6	10.75	8.2	9.97	7.29	7.74	8.68	*	*	-	-

76 / 157

[0900]

[表429]

2591

2592

NT2RP2003125	148.62	154.21	165.76	186.93	87.76	119.3	202.39	170.18	181.03				
NT2RP2003127	8.19	13.35	12.24	9.29	4.08	4.24	7.52	6.27	6.75				
NT2RP2003128	5.37	4.98	2.74	4.8	1.83	1.5	2.73	2.54	2.23				
NT2RP2003127	9.57	8.81	8.47	10.46	3.57	7.13	10.02	8.73	11.39				
NT2RP2003134	289.53	193.02	256.72	147.84	144.9	131.2	108.58	156.88	150.94	*	*	-	-
NT2RP2003144	1.24	1.09	7.94	3.13	0.31	1.29	1.85	1.69	1.77				
NT2RP2003144	8.05	4.25	18.47	5.03	5.56	4.21	3.03	3.24	3.73				
NT2RP2003150	10.24	5.68	13.18	8.31	3.11	3.88	5.08	3.43	3.42				
NT2RP2003167	3.48	4.11	8.05	5.53	2.9	2.53	3.08	2.08	2.29				
NT2RP2003168	154.65	121	143.84	65.15	86.73	95.34	71.82	47.48	71.44	*	**	-	-
NT2RP2003181	7.49	10.19	13.18	11.02	5.25	6.57	6.09	5.72	4.72	*		-	-
NT2RP2003164	3.55	3.69	5.64	7.04	3.38	4.04	7.48	6.26	5.34				
NT2RP2003185	12.93	8.52	16.52	10.68	5.85	8.09	3.19	7.18	6.13				
NT2RP2003177	20.41	17.38	32.83	19.51	10.51	7.06	22.73	21.74	21.1				
NT2RP2003179	13.75	8.83	17.7	6.83	5.73	5.2	6.82	13.09	8.84	*		-	-
NT2RP2003194	60.38	55.91	80.01	51.34	60.6	65.85	42.59	38.48	56.22				
NT2RP2003206	4.24	5.6	7.88	6.81	4.67	3.88	3.76	3.9	1.77				
NT2RP2003210	9.9	9.38	18.05	7.55	5.04	4.08	4.47	4.65	4.08	*	**	-	-
NT2RP2003227	13.08	17.16	19.18	9.3	7.47	5.3	8.38	6.37	6.85	*	**	-	-
NT2RP2003228	16.97	20.02	17.73	19.68	23.22	23.38	16.98	9.41	24.11				
NT2RP2003230	8.37	4.28	11.24	9.26	4.24	4.38	5.34	6.55	20.33				
NT2RP2003231	19.49	17.75	28.22	18.04	13.83	13.5	4.32	15.98	15.62				
NT2RP2003237	7.53	6.29	14.94	7.53	4.21	4.44	2.72	5.12	9.6				
NT2RP2003235	14.69	12.62	13.82	13.84	7.63	7.59	6.01	11.3	7.74	*		-	-
NT2RP2003243	18.5	16.09	21.39	14.68	13.35	11.44	13.72	14.6	10.6	*	*	-	-
NT2RP2003265	4.18	3.24	4.88	3.11	1.7	1.78	2.29	2.88	1.56	*	*	-	-
NT2RP2003267	60.51	57	71.61	43.05	41.52	45.25	62.67	46.27	56.1	*		-	-
NT2RP2003272	18.93	12.31	21.18	15.82	7.25	9.68	9.71	5.82	7.07	*		-	-
NT2RP2003277	51.75	42.05	60.1	30.74	24.84	22.69	27.09	34.52	27.77	*	*	-	-
NT2RP2003283	28.6	31.79	38.98	27.18	28.4	22.37	7.26	22.23	17.9	*	*	-	-
NT2RP2003285	5.27	5.54	11.77	5.48	3.75	3.02	2.08	4.89	2.65				
NT2RP2003293	449.82	544.88	531.61	480.84	613.42	591.27	492.68	295.17	642.99				
NT2RP2003295	31.05	28.41	29.77	21.7	14.01	19.1	19.18	18.48	19.51	**	**	-	-
NT2RP2003297	16.92	17.67	19.83	9.38	11.73	7.84	3.24	6.25	6.08	**	**	-	-
NT2RP2003300	2.42	4.38	4.19	6.32	2.8	1.68	2.33	1.69	1.31				
NT2RP2003302	4.27	4.34	3.48	7.02	4.29	2.92	3.66	4.67	4.2				
NT2RP2003307	17.52	15.84	28.87	14.93	9.5	10.4	16.58	17.1	22.41				
NT2RP2003308	7.26	8.68	14.15	6.9	4.51	4.44	3.22	4.78	6.65				
NT2RP2003311	29.64	20.85	41.84	22.79	16.59	15.7	26.2	28.58	26.19				
NT2RP2003329	11.38	11.27	19.3	9.84	10.72	8.3	6.72	6.84	5.55	*		-	-
NT2RP2003339	44.08	51.39	68.68	63.28	61.45	41.82	35.03	31.79	30.76	*		-	-
NT2RP2003345	25.41	28.87	30.12	15.65	24.89	17.7	13.82	14.02	13.58	*	**	-	-
NT2RP2003347	8.25	20.81	11.17	7.76	7.09	4.08	5.16	5.37	4.53				
NT2RP2003367	15.71	9.07	15.82	25.01	12.88	13.87	19.6	13.65	17.34				
NT2RP2003369	28.62	21.24	26	8.88	11.95	10.09	4.71	10.13	9.11	**	**	-	-
NT2RP2003383	14.02	11.63	23.31	10.33	7.37	5.29	4.94	12.01	8.63				
NT2RP2003390	5.02	4.43	9.92	6.51	4.16	1.23	2.19	2.28	1.48				
NT2RP2003391	7.9	10.18	14.85	7.88	4.63	3.15	3.6	3.53	4.69	*		-	-
NT2RP2003393	17.88	10.78	13.47	6.59	12.87	3.87	3.38	3.46	4.48	**		-	-
NT2RP2003394	86	76.58	98.35	85.98	75.14	89.1	44.7	65.38	51.85	**		-	-
NT2RP2003401	6.44	4.6	7.19	7.26	3.17	2.2	2.64	2.79	2.35	*		-	-
NT2RP2003403	18.73	19.58	18.59	18.65	31.07	15.94	9.34	7.98	7.02	**		-	-
NT2RP2003433	8.13	7.84	13.82	4.56	2.38	2.9	0.77	3.05	2.35	*	*	-	-
NT2RP2003445	11.02	10.94	17.55	8.15	5.02	3.19	3.72	5.83	4.71	*	*	-	-
NT2RP2003446	7.38	5.87	15.32	8.9	3.31	1.64	2.84	3.34	2.51				
NT2RP2003458	14.58	16.09	17.24	7.76	9.78	4.68	6.41	6.81	4.84	**	**	-	-
NT2RP2003468	163.69	136.64	181.26	110.44	140.48	151.43	83.63	87.04	121.38	*		-	-
NT2RP2003468	5.44	3.7	6.03	4.74	2.31	1.35	2.28	1.42	1.82	**		-	-
NT2RP2003470	4.63	5.55	5.49	5.65	3.19	3.4	4.42	1.43	3				
NT2RP2003471	12.29	11.22	13.81	11.17	23.08	8.88	5.31	3.9	2.98	**		-	-
NT2RP2003480	133.45	297.88	151.87	131.22	166.23	117.7	104.84	202.75	202.47				
NT2RP2003495	2.81	3.4	10.31	5.19	1.55	2.41	0.68	1.48	0.7				
NT2RP2003499	5.25	3.3	9.47	3.94	2.15	2.35	2.44	6.25	5.89				
NT2RP2003505	1.65	1.94	6.72	4.41	2.48	0.11	0.21	3	0				

77 / 157

2593

2594

NT2RP2003508	18.91	15.11	21.41	10.84	13.4	7.04	4.15	4.83	3.94	*	**	-	-
NT2RP2003511	12.05	3.9	4.84	8.42	8.48	3	2.22	2.49	1.23			-	-
NT2RP2003513	5.02	4.48	4.09	4.53	1.84	2.04	2.58	1.65	3.7	*		-	-
NT2RP2003517	3.09	1.94	2.24	4.01	1.27	1.81	1.81	0.62	1.53			-	-
NT2RP2003522	5.79	5.52	8.11	4.81	3.87	1.78	1.31	4.88	5.44			-	-
NT2RP2003526	12.34	13.08	18.44	6.02	5.34	4.03	6.04	8.8	5.02	**	**	-	-
NT2RP2003533	89.01	187.21	169.95	95.54	157.31	89.82	178.82	120.18	151.89			-	-
NT2RP2003541	3.87	3.34	5.25	3.35	1.69	1.28	1.11	0.78	0.47	**		-	-
NT2RP2003543	6.02	7.31	6.9	4.81	3.89	1.42	3.96	1.71	3.08	*	**	-	-
NT2RP2003545	16.66	18.91	37	10.57	5.83	7.38	9	7.02	10.57	*	*	-	-
NT2RP2003559	24.73	30.59	37.42	18.05	13.28	13.04	17.81	5.8	8.19	*	*	-	-
NT2RP2003564	4.21	3.5	2.84	5.11	2.21	2.78	1.47	1.11	1.54	**		-	-
NT2RP2003565	5.58	8.05	9.75	8.3	2.97	4.12	3.13	4.62	3.34			-	-
NT2RP2003567	52.47	49.81	31.71	25.62	15.66	18.12	8.17	10.23	14.83	*	*	-	-
NT2RP2003575	228.18	422	284.34	79.35	234.74	114.07	248.45	399.29	319.2			-	-
NT2RP2003578	12.51	13.48	14.93	11.41	10.85	7.47	9.35	12.38	11.68	*		-	-
NT2RP2003579	6.35	14.72	8.69	5.77	6	3.85	5.14	3.96	3.9			-	-
NT2RP2003581	5.18	8.39	5.74	5.34	1.72	1.57	1.66	0.89	1.73	**		-	-
NT2RP2003587	5.23	4.31	3.93	4.85	1.48	2.81	2.91	3.12	2.6	*		-	-
NT2RP2003590	10.26	7.56	7.31	8.22	4.4	10.25	8.81	5.02	6.24			-	-
NT2RP2003593	30.36	28.83	29	16.89	17.04	17.88	10.43	21.71	16.36	**	*	-	-
NT2RP2003596	9.32	8.79	11.25	13.55	9.73	9.49	8.79	14.74	9.47			-	-
NT2RP2003599	28.27	13.17	35	8.05	27.81	6.94	3.86	4	10.63			-	-
NT2RP2003600	19.88	30.4	31.86	22.5	9.14	19.7	15.99	13.53	9.08	*		-	-
NT2RP2003604	6.88	9.89	7.47	5.4	1.89	1.85	3.39	3.16	2.76	*	*	-	-
NT2RP2003629	8.19	7.7	5.28	5.58	3.64	2.51	3.1	3.51	2.87	*		-	-
NT2RP2003630	4.6	3.1	1.8	4.67	0.47	1.48	1.48	1.9	2.02			-	-
NT2RP2003643	14.28	14.44	12.75	14.01	7.67	9.12	19.42	10.04	13.07			-	-
NT2RP2003655	7.62	5.24	7.73	7.33	2.44	6.8	2.97	5.19	4.67	*		-	-
NT2RP2003664	8.98	12.67	11.03	11.29	17.02	7.98	3.52	3.83	3.73	**		-	-
NT2RP2003688	6.83	2.7	3.15	6.14	1.92	4.81	2.43	9.15	4.25			-	-
NT2RP2003687	2.45	2.75	3.32	4.63	0.64	2.77	0.71	2.3	0.81			-	-
NT2RP2003691	2.88	0.41	1.55	4.29	1.06	2.07	1.89	2.35	1.61			-	-
NT2RP2003702	7.99	4.44	8.03	9.77	1.89	6.58	5.28	5.52	3.42			-	-
NT2RP2003704	6.11	9.87	8.13	7.33	6.55	3.79	6.43	2.95	5.24			-	-
NT2RP2003706	7.36	9.13	9.09	6.95	3.88	3.8	6.16	2.16	4.58	*	*	-	-
NT2RP2003713	10.55	5.71	14.4	8.76	4.92	4.56	5.34	3.61	4.08			-	-
NT2RP2003714	8.27	4.37	12.83	8.63	2.38	6.34	8.02	5.56	6.23			-	-
NT2RP2003727	13.72	9.91	16.85	14.5	8.49	5.28	4.4	3.92	5.54	*		-	-
NT2RP2003737	5.45	5.17	12.85	7.78	3.42	3.25	3.61	5.76	7.42			-	-
NT2RP2003751	4.58	1.12	4.62	4.58	0.42	0.71	2.16	1.87	0.69			-	-
NT2RP2003780	5.71	6.37	7.89	7.18	2.44	3.65	5.58	4.39	4.01			-	-
NT2RP2003784	4.38	3.82	3.93	5.42	1.13	2.52	3.39	1.74	2.45	*		-	-
NT2RP2003789	20.81	10.06	18.85	18.88	9.37	7.83	5.73	10.32	7.17			-	-
NT2RP2003770	48.73	23.49	36.27	27.58	17.68	15.16	13.63	17.82	25.92			-	-
NT2RP2003777	15.17	10.4	15.63	9.4	5.35	5.3	5.51	8.22	8.17	*	*	-	-
NT2RP2003781	85.09	53.2	80.23	44.9	32.89	44.6	88.34	75.84	76.38	*		-	-
NT2RP2003785	11.82	18.04	20.61	13.78	15.2	6.86	6.32	9.84	5.9	*		-	-
NT2RP2003793	16.4	16.81	28.84	17.84	16.4	19.5	8.6	5.88	10.68	*		-	-
NT2RP2003806	22.61	30.87	30.72	19.98	15.24	17.95	21.46	20.85	31.8	*		-	-
NT2RP2003825	121.72	69.01	127.35	108.65	103.08	122.02	108.88	101.92	114.28			-	-
NT2RP2003840	11.68	9.2	14.49	8.94	8.01	4.99	5.08	8.29	6.32	*		-	-
NT2RP2003857	12.37	17.33	20.12	13.2	12.38	5.39	3.92	13.14	19.37			-	-
NT2RP2003859	19.27	8.66	21.11	12.54	7.05	8.25	8.14	9.73	13.41			-	-
NT2RP2003871	24.21	21.74	19.35	11.95	9.8	10.67	5.81	8.41	8.75	**	**	-	-
NT2RP2003876	10.63	11.38	18.68	19.02	25.1	11.05	7.58	10.36	8.8			-	-
NT2RP2003878	8.32	9.1	10.32	8.36	3.78	3.6	4.02	3.95	3.32	**		-	-
NT2RP2003885	6.08	8.46	6.77	5.72	4.33	2.2	4.11	3.32	4.21	*		-	-
NT2RP2003898	18.47	22.78	20.73	12.55	8.77	11.94	7.99	8.27	7.33	*	**	-	-
NT2RP2003902	21.89	10.54	20.35	18.69	8	10.68	8.28	15.68	16.89			-	-
NT2RP2003912	35.18	73.65	32.26	21.86	23.82	12.49	12.87	18.46	19.33			-	-
NT2RP2003931	18.31	15.84	40.63	16.98	12.81	14.09	15.19	20.43	45.95			-	-
NT2RP2003940	28.29	28.19	28.81	19.73	11.84	14	20.12	28.2	22.68	**		-	-
NT2RP2003950	10.42	7.47	13.08	10.4	5.75	5.62	8.83	8.37	6.11			-	-

78 / 157

2595

2596

NT2RP2003952	8.61	6.4	10.47	5.99	3.42	3.14	1.11	4	4.05	*	*	-	-
NT2RP2003968	14.45	22.39	24.81	11.6	25.12	6.38	6.57	9.01	14.39			-	-
NT2RP2003976	33.5	38.02	32.11	28.15	26.29	22.25	25.49	23.11	18.45	*	**	-	-
NT2RP2003981	10.38	7.73	15.47	7.98	3.66	7.9	3.93	9.3	6.45			-	-
NT2RP2003984	21.86	17.33	21.82	17.29	10	7.7	4.79	10.99	18.69			-	-
NT2RP2003986	29.04	20.89	42.5	17.22	10.08	14.95	20.31	27.16	19.47			-	-
NT2RP2003988	17.98	13.51	22.98	15.07	9.71	10.14	10.31	12.25	15.9			-	-
NT2RP2004011	78.9	66.49	88.74	48.94	44.67	24.09	28.41	25.31	32.75	**	**	-	-
NT2RP2004014	12.05	10.78	9.85	7.34	4.69	5.32	4.75	6.68	7.08	**	**	-	-
NT2RP2004036	38.11	34.32	48.47	33.87	30.52	34.54	23.89	26.94	32.83			-	-
NT2RP2004041	15.68	9.6	13.65	12.24	6.88	6.29	9.48	7.39	4.47			-	-
NT2RP2004042	10.47	6.9	14.24	8.29	4.18	4.2	3.12	6.29	6.45			-	-
NT2RP2004049	50.08	29.35	58.88	54.03	47.21	47.08	23.48	30.4	35.04			-	-
NT2RP2004080	14.43	11.83	19.24	13.07	7.66	8.82	8.25	9.01	6.7	*		-	-
NT2RP2004085	9.22	7.85	15.12	7.14	3.11	3.37	2.62	3.61	3.15	*		-	-
NT2RP2004089	13.07	11	10.34	9.29	5.69	3.82	4.85	6.14	4.78	*		-	-
NT2RP2004076	7.23	8.97	8.94	6.15	5.2	2.62	3.55	3.66	4.01	*	**	-	-
NT2RP2004080	16.76	19.32	21.22	10.81	7.94	4.93	7.71	6.49	6.54	**	**	-	-
NT2RP2004081	12.13	9.06	12.85	7.67	4.89	5.2	4.78	5.28	3.41	*	**	-	-
NT2RP2004088	14.04	9.1	18.6	7.17	4.65	6.94	5.74	6.96	7.08			-	-
NT2RP2004108	15	13.48	22.62	10.08	6.21	6.99	5.4	7.73	10.17	*	*	-	-
NT2RP2004124	8.21	9.75	17.19	8.68	5.64	4.42	4.13	6.1	2.83			-	-
NT2RP2004130	14.64	18.18	18.71	14.91	10.54	10.38	6.64	6.22	8.03	**		-	-
NT2RP2004133	9.17	7.74	14.55	8.02	4.47	1.68	3.8	2.83	1.86	*		-	-
NT2RP2004141	13.91	6.7	8.85	9.25	5.97	4.12	2.68	2.87	3.2	*		-	-
NT2RP2004142	20.24	20.68	23.15	12.58	8.99	0.11	7.78	8.1	8.83	*	**	-	-
NT2RP2004152	12.94	17.34	28.22	13.04	12.79	7.85	8.66	8.13	4.02			-	-
NT2RP2004185	23.06	20.1	27.11	14.41	8.19	16.16	10.81	12.82	15.14	*	*	-	-
NT2RP2004170	14.84	22.1	21.27	11.58	5.42	12.62	7.64	14.97	10.1	*		-	-
NT2RP2004172	11.82	8.43	19.13	9.78	6.86	6.4	5.08	5.40	3.29			-	-
NT2RP2004175	18.04	24.68	27.17	14.08	13.85	12.56	15.42	21.18	13.53			-	-
NT2RP2004179	11.06	11.88	18.99	6.12	3.68	3.62	3.12	4.46	3.04	*	*	-	-
NT2RP2004187	10.62	14.3	17.06	10.83	6.49	6	6.3	3.3	5.65	*		-	-
NT2RP2004180	11.44	13.08	11.65	11.65	16.98	6.89	14.27	4.91	8.09			-	-
NT2RP2004184	34.23	32.85	35.34	19.98	14.64	15.16	22.5	12.25	14.92	**	**	-	-
NT2RP2004186	30.83	39.33	27.33	12.84	5.98	11.23	10.98	13.1	14.08	**	**	-	-
NT2RP2004205	29.01	29.84	37.62	20.1	10.4	16.74	17.25	16.98	24.08	*	*	-	-
NT2RP2004207	8.99	13.88	16.49	8.05	7.02	4.72	8.6	7.78	25.16			-	-
NT2RP2004226	17.68	11.75	16.87	11.26	9.8	6.13	9.93	6.52	8.04	*		-	-
NT2RP2004232	9.38	7.39	12.33	6.45	7.37	3.57	6.49	3.42	8.98			-	-
NT2RP2004239	9.12	12.33	10.22	7.52	5.19	3.65	2.71	5.07	3.29	*	**	-	-
NT2RP2004240	34.87	45.84	33.68	20.39	18.64	8.67	14.35	8.97	4.81	*	**	-	-
NT2RP2004242	15.84	15.83	16.65	14.76	9.29	9.76	10.22	6.01	6.98	**		-	-
NT2RP2004245	7.49	8.21	10.79	7.33	2.04	4.45	4.31	4.59	4.95	*		-	-
NT2RP2004270	62.73	47.21	68.41	48.4	33.04	45.75	50.52	75.49	70.49			-	-
NT2RP2004300	12.51	8.53	13.1	8.54	9.07	5.04	6.79	9.38	7.79			-	-
NT2RP2004304	18.04	25.81	28.38	10.74	8.89	8.88	7.57	8.7	9.4	*	*	-	-
NT2RP2004313	5.32	5.95	8.95	4.78	1.11	1.27	2.88	2.46	2.35	*		-	-
NT2RP2004316	7.59	19.66	19.22	7.37	4.16	2.63	3.4	2.3	5.68	*		-	-
NT2RP2004321	11.35	10.01	12.72	6.18	1.74	2.16	3.62	2.37	4.34	**	**	-	-
NT2RP2004336	6.56	6.01	5.01	7.15	3.83	3.55	3.88	1.85	3.84	*		-	-
NT2RP2004339	88.87	54.65	84.87	50.21	27.02	70.76	73.63	111.22	98.82			-	-
NT2RP2004347	9.68	12.29	15.81	9.78	3.94	5.75	4.47	6.18	9.27			-	-
NT2RP2004364	17.82	15.18	19.5	11.98	7.62	12.3	9.63	15.21	13.78	*		-	-
NT2RP2004365	16.43	15.15	23.6	8.87	8.61	4.88	6.23	5.81	4.59	**	**	-	-
NT2RP2004368	6.71	10.68	9.69	7.66	3.53	5.17	5.46	3.22	3.29	*		-	-
NT2RP2004373	12.26	11.21	11.74	7.28	2.71	4.58	3.35	3.56	4.51	**	**	-	-
NT2RP2004376	69.47	77.52	81.72	36.98	16.48	21.69	34.48	29.81	18.56	**	**	-	-
NT2RP2004389	11.31	16.38	12.26	9.91	2.21	4.95	5.82	4.38	4.02	*	**	-	-
NT2RP2004392	67.13	31.73	43.78	20.38	16.25	19.24	10.85	16.27	14.58	*	*	-	-
NT2RP2004396	13.42	9.2	11.08	9.69	3.89	7.39	5.87	7.43	7.78	*		-	-
NT2RP2004399	25.77	19.74	26.11	15.91	12.58	15.11	9.21	12.21	10.01	*	**	-	-
NT2RP2004400	10.29	9.25	9.33	9.9	4.17	8.44	7.31	8.61	9.43			-	-
NT2RP2004404	142.91	126.47	123.47	87.01	85.43	109.28	171.94	127.81	174.94	*		-	-

2597

2598

NT2RP2004410	39.53	65.74	46.52	39.11	18.74	18.35	21.28	23.23	25.45	*	-
NT2RP2004412	14.74	10.38	20.72	26.06	1.98	8.17	8.89	4.91	6.36	*	-
NT2RP2004414	8.28	7	7.21	7.75	3.36	2.92	8.71	6.3	5.47	*	-
NT2RP2004425	10.2	6.28	18.09	7.43	2.9	4.39	7.1	5.14	4.24	*	-
NT2RP2004447	7.9	4.2	15.8	7.23	1.76	4.51	4.43	8.04	8.17	*	-
NT2RP2004463	53.44	48.94	45.23	51.07	39.96	24.58	38.74	19.02	17.11	*	-
NT2RP2004476	18.45	12.59	23.94	14.93	16.73	11.7	18.21	16.49	14.59	*	-
NT2RP2004488	10.21	10.37	13.58	8.53	5.46	7.65	8.97	6.18	4.15	*	-
NT2RP2004490	6.68	9.48	11.96	10.41	4.06	3.06	6.4	5.15	3.37	*	-
NT2RP2004495	59.84	40.88	52.11	47.08	29.44	39.52	53.9	45.74	40.44	*	-
NT2RP2004512	17.42	8.07	18.38	11.03	4.08	6.19	1.75	5.77	3.87	*	-
NT2RP2004523	26.61	22.99	26.09	20.65	9.58	18.73	12.96	22.93	30.09	*	-
NT2RP2004524	21.22	18.85	29.35	17.14	12.55	10.77	11.27	10.59	14.51	*	-
NT2RP2004535	39.58	17.04	37.57	26.29	26.05	14.38	14.91	28.85	19.32	*	-
NT2RP2004538	145.51	113.79	134.59	95.27	89.81	81.41	118.33	103.7	122.39	*	-
NT2RP2004548	21.33	25.87	20.88	16.6	11.36	9.17	14.62	15.2	15	*	-
NT2RP2004551	8.55	11.05	18.95	8.38	5.83	4.59	5.77	6.04	8.54	*	-
NT2RP2004556	187.44	116.09	115.54	100.64	63.01	68.21	138.61	56.34	79.2	*	-
NT2RP2004568	25.54	15.99	32.04	18.77	7.42	18.63	9.21	19.06	44.48	*	-
NT2RP2004580	28.48	22.84	30.45	23.04	12.01	15.1	24.34	30.83	24.49	*	-
NT2RP2004585	104.19	75.97	102.59	102.16	96.71	82.97	64.51	65.78	69.08	*	-
NT2RP2004587	7.64	7.71	10.73	8.38	2.94	4.03	2.08	3.86	11.64	*	-
NT2RP2004594	9.49	13.7	13.53	10.84	10.06	5.98	7.18	7.17	14.25	*	-
NT2RP2004600	11.79	11.85	12.42	9.8	6.39	8.69	9.84	7.14	9.81	*	-
NT2RP2004602	16.85	24.33	21.89	13.29	12.48	8.02	12.27	7.59	7.04	*	-
NT2RP2004606	282.85	262.35	268.41	181.43	249.55	245.28	257.06	189.04	158.13	*	-
NT2RP2004614	9.31	7.04	11.05	6.07	2.87	4.48	3.52	3.61	6.4	*	-
NT2RP2004648	12.5	37.87	41.82	14.38	8.31	11.65	10.13	8.24	22.69	*	-
NT2RP2004655	27.02	40.5	41.28	26.01	23.59	13	4.9	10	13.87	**	-
NT2RP2004664	12.3	12.14	15.05	9.84	6.17	5.52	2.65	4.39	2.54	*	-
NT2RP2004670	9.81	11.79	9.6	6.17	5.85	2.45	3.26	4.14	2.42	*	-
NT2RP2004675	22.26	31.4	25.45	19.55	10.99	11.78	17.4	17	31.58	*	-
NT2RP2004681	11.82	17.49	14.89	8.26	5.82	3.58	5.62	7.61	5.68	*	-
NT2RP2004689	5.34	25.14	16.38	16.38	9.9	10.44	10.84	15.98	14.64	*	-
NT2RP2004709	38.28	22.19	35.75	22.32	11.28	18.42	7.39	20.31	16.27	*	-
NT2RP2004710	23.78	32.41	33.69	19.19	16.47	14	10.16	17.13	21.44	*	-
NT2RP2004721	15.3	9.83	18.51	7.84	2.48	4.31	3.84	6.3	2.89	*	-
NT2RP2004735	12.51	12.2	14.59	12.94	8.12	6.27	5.93	10.35	6.34	*	-
NT2RP2004743	18.88	18.19	16.11	11.52	9.54	6.56	6.31	8.72	9.81	**	-
NT2RP2004750	59.2	58.12	63.52	47.5	43.74	38.18	59.22	63.07	62.81	**	-
NT2RP2004755	27.3	33.28	34.09	17.59	21.76	19.12	17.05	12.23	8.96	**	-
NT2RP2004767	18.81	19.28	18.07	14.81	10.15	10.32	11.83	8.67	9.89	*	-
NT2RP2004768	39.27	17.22	26.44	49.02	23.23	30.46	23.63	20.93	17.78	*	-
NT2RP2004775	10.4	6.35	15.81	7.59	3.37	4.94	1.86	3.35	3.03	*	-
NT2RP2004791	35.3	37.93	41.25	33.85	20.1	18.04	17.4	29.11	27.1	*	-
NT2RP2004794	64.45	52.25	59.58	37.23	24.39	18.04	14.21	17.04	17.83	*	-
NT2RP2004795	16.82	13.04	16.28	12.97	8.62	3.98	5.33	6.36	8.84	**	-
NT2RP2004799	18.76	14.17	14.52	9.03	7.47	5.53	2.73	2.76	3.58	**	-
NT2RP2004802	42.99	46.43	41.15	46.78	39.32	30.76	28.59	29.92	38.28	*	-
NT2RP2004810	18.3	18.68	17.35	9.03	6.57	4.19	8.85	4.21	4.39	**	-
NT2RP2004818	39.61	34.82	49.2	16.85	15.88	3.1	5.7	11.84	0.57	*	-
NT2RP2004837	65.01	32.88	59.21	31.78	19.92	27.08	12.99	11.53	9.41	*	-
NT2RP2004841	12.28	19.6	17.04	13.83	8.03	5.68	4.26	5.92	8.44	*	-
NT2RP2004847	58	45.24	59.59	31.54	30.59	31.44	43.02	39.63	29.88	**	-
NT2RP2004881	8.61	8.78	10.74	6.21	2.52	2.33	3.42	5.23	3.33	*	-
NT2RP2004897	8.08	8.52	10.02	7.86	4.8	3.34	2.33	3.07	4.68	**	-
NT2RP2004932	11.95	10.48	13.81	9.34	4.73	6.48	4.87	3.86	5	*	-
NT2RP2004933	6.78	7.42	14.62	6.21	3.1	2.23	3.41	1.97	1.54	*	-
NT2RP2004936	9.66	18.67	21.72	10.34	4.39	7.29	2.62	3.9	3.25	*	-
NT2RP2004951	8.57	7.11	16.72	9.28	2.32	5.42	3.11	6.13	4.04	*	-
NT2RP2004958	12.5	13.69	22.18	6.53	5.09	6.63	3.92	5.67	5.43	*	-
NT2RP2004961	15.97	14.84	23.99	12.05	7.35	8.71	10.28	8.72	8.04	*	-
NT2RP2004962	15.99	15.58	20.13	13.9	10.21	9.93	9.78	10.65	9.63	*	-
NT2RP2004986	6.94	8.72	7.55	5.31	3.88	2.7	4.34	2.81	2.83	*	-

80 / 157

2599

2600

NT2RP2004987	18.8	14.08	15.99	13.12	6.04	9.98	11.43	12.91	9.85	*	-
NT2RP2004974	8.02	8.78	7.74	5.74	3.27	3.73	5.44	2.35	2.1	**	-
NT2RP2004978	14.12	27.31	13.7	11.98	4.65	6.79	4.18	9.57	4.35	*	-
NT2RP2004982	4.88	6.54	10.22	4.28	1.48	0.65	1.99	1	1.83	*	-
NT2RP2004985	209.85	195.75	215.75	188.58	153.94	208.41	168.37	224.84	192.82	*	-
NT2RP2004989	18.04	12.97	26.42	17.44	8.41	10.32	8.65	18.31	9.11	*	-
NT2RP2005000	9.18	10.25	11.07	8.89	2.87	3.28	5.48	4.95	3.42	*	**
NT2RP2005001	18.08	20.79	40.08	15.62	11.69	12.08	12.25	15.19	13.27	*	-
NT2RP2005003	15.88	14.42	15.05	11.32	5.53	6.52	11.83	11.19	8.75	*	-
NT2RP2005012	40.61	67.68	43.81	20.05	11.76	18.65	21.02	13.16	20.53	**	**
NT2RP2005018	6.29	10.32	12.32	7.2	3.34	4.91	3.54	5.22	4.18	*	-
NT2RP2005020	138.98	109.02	105.29	97.72	59.81	79.64	117.66	176.16	138.64	*	-
NT2RP2005022	8.77	8.01	12.34	8.19	7.74	5.07	4.95	7.18	9.74	*	-
NT2RP2005027	53.48	62.9	53.33	58.43	50.24	56.5	33.74	38.11	31.82	**	-
NT2RP2005031	6.85	8.81	9.58	5.73	4.02	3.35	3.46	3.48	3.14	*	**
NT2RP2005037	16.82	18.46	19.47	6.47	4.35	2.93	2.65	3.66	8.14	**	**
NT2RP2005038	6.96	8.88	11.09	7.2	2.46	3.49	4.34	2.45	1.1	*	-
NT2RP2005048	39.21	28.46	29.83	31.24	22.21	43.14	32.34	48.11	37.29	*	-
NT2RP2005049	79.3	47.14	64.37	36.64	18.08	20.34	23.98	33.08	29.72	*	-
NT2RP2005073	67.78	64.88	62.64	70.16	48.08	81.7	55.95	55.76	57.19	*	-
NT2RP2005097	10.25	7.51	14.69	8.49	1.21	4.7	3.76	2.8	2.35	*	-
NT2RP2005108	13.67	11.69	13.11	8.87	5.69	6.68	5.31	8.56	5.73	**	**
NT2RP2005116	11.75	10.87	12.14	10.24	4.17	4.29	7.43	5.88	6.2	**	-
NT2RP2005126	21.43	28.84	28.25	14.77	6.89	5.64	7.77	5.68	5.93	*	**
NT2RP2005135	10.21	11.92	11.29	12.06	8.02	7.4	9.29	7.72	6.92	*	-
NT2RP2005139	4.47	5.9	6.41	5.94	3.02	9.46	4.16	4.52	4.09	*	-
NT2RP2005140	10.93	7.52	7.44	7.58	3.55	5.44	3	4.22	2.28	*	-
NT2RP2005144	13.73	10.64	13.89	8.29	4.49	7.83	1.84	3.98	2.4	*	**
NT2RP2005147	22.68	15.06	21.74	11.07	5.61	11.36	5.53	13.64	8.82	*	-
NT2RP2005148	19.34	14.88	12.3	16.32	7.6	14.47	18.68	18.99	24	*	-
NT2RP2005159	9.62	7.06	9.83	13.34	5.14	7.82	4.38	3.58	6.05	*	-
NT2RP2005162	14.64	12.6	16.31	14.53	7.84	8	8.72	8.34	7.02	**	-
NT2RP2005181	25.6	37.52	33.86	15.82	13.92	18.66	24.95	23.28	21.32	*	-
NT2RP2005188	10.58	6.62	14.16	11.05	5.89	4.43	4.83	6.74	4.38	*	-
NT2RP2005181	4.05	2.99	8.75	6.74	2.11	1.29	1.74	3.19	2.03	*	-
NT2RP2005204	14.87	10.24	20.73	15.12	7.06	10.51	11.01	8.88	10.24	*	-
NT2RP2005219	24.24	18.16	32.19	28.01	12.29	15.28	21.03	18.69	23.06	*	-
NT2RP2005227	12.24	9.75	17.81	17.67	5.27	7.45	11.51	12.08	11.25	*	-
NT2RP2005237	103.35	89.54	108.29	128.64	78.44	98.83	81.16	75.94	83.42	*	-
NT2RP2005239	4.88	8.22	9.62	8.57	2.23	3.55	3.5	4.01	4.35	*	-
NT2RP2005247	22.45	26.01	32.27	32.43	14.24	19.04	34.79	31.98	33.52	*	-
NT2RP2005254	36.5	15.86	25.1	17.04	11.38	15.24	5.51	13.65	13.69	*	-
NT2RP2005270	25.5	13.21	29.5	13.51	9.93	13.53	11.24	14.21	22.67	*	-
NT2RP2005276	40.5	21.43	40.17	28.18	14.43	18.68	10.58	8.43	8.36	*	-
NT2RP2005287	16.9	10.22	25.14	13.6	9.12	7.29	8.33	8.76	8.8	*	-
NT2RP2005288	8.35	8.09	8.85	7.18	3.2	3.43	4.54	4.39	3.3	*	-
NT2RP2005289	19.76	15.26	21.43	14.2	18.67	10.92	15.53	14.93	15.73	*	-
NT2RP2005293	9.78	11.07	10.85	8.57	5.47	3.14	6.98	5.79	5.74	*	**
NT2RP2005315	16.84	21	21.65	20.87	8.72	12.4	13.48	9.21	11.44	*	-
NT2RP2005322	33.55	31.7	65.28	25.82	25.53	28.43	14.38	22.11	31.46	*	-
NT2RP2005325	54.2	44	55.99	38.68	30.92	29.07	18.94	27.07	30.65	**	**
NT2RP2005336	28.38	12.98	27.98	11.35	7.87	9.01	10.99	13.9	14.17	*	-
NT2RP2005343	12.61	22.98	14.15	9.9	5.77	8.7	7.81	10.19	10.12	*	-
NT2RP2005344	6.04	8.8	10.41	6.82	3.18	3.71	4.76	3.88	3.88	*	-
NT2RP2005347	7.59	13.3	8.89	7.33	4.35	4.03	4.9	3.27	3.77	*	-
NT2RP2005354	41.58	40.59	43.7	31.3	21.57	29.41	26.2	24.6	22.64	**	**
NT2RP2005358	43.39	32.77	43.03	81.08	32.41	50.49	63.9	54.2	68.64	*	+
NT2RP2005360	11.79	10	18.74	11.01	6.71	7.24	3.72	6.21	9.78	*	-
NT2RP2005378	26.24	21.48	29.8	11.99	9.12	7.48	6.73	9.68	15.33	**	*
NT2RP2005391	8.87	18.64	25.48	15.21	9.78	9	4.05	7.14	6.04	*	-
NT2RP2005393	18.68	11.72	17.99	12.44	7.52	10.75	12.81	18.43	12.18	*	-
NT2RP2005407	7.79	10.99	12.03	9.14	5.78	6.4	4.56	6.12	6.26	*	-
NT2RP2005419	14.03	13.17	13.95	13.78	13.3	12.37	11.42	8.28	10.02	**	-
NT2RP2005425	22.13	23.63	26.77	16.49	17.88	16.5	26.07	23.18	23.96	*	-

81 / 157

2601

2602

NT2RP2005429	20.77	21.25	24.75	28.87	19.8	22.9	31.5	41.11	27.5				
NT2RP2005436	21.72	21.01	38.27	15.82	10.49	11.17	6.19	10.76	15.31				
NT2RP2005441	12.34	10.42	16.43	6.82	3.95	2.88	3.88	5.58	7.88	0	0	-	-
NT2RP2005442	100.99	76.34	97.05	85.22	107.34	84.16	56.06	95.44	78.81				
NT2RP2005444	44.56	42.84	50.52	45	41.48	37.39	32.32	34.33	40.3	0	0	-	-
NT2RP2005453	6.3	5.27	7.3	6.26	2.81	2.81	2.27	2.79	2.65	00	00	-	-
NT2RP2005457	85.08	108.4	103.3	51.25	78.76	69.99	40.37	68.9	47.82	0	0	-	-
NT2RP2005458	19.83	18.76	13.48	12.24	12.61	6.33	8.28	12.39	11.01	0	0	-	-
NT2RP2005463	26.71	23.66	25.36	21.35	13.39	12.75	11.08	11.87	8.93	0	00	-	-
NT2RP2005464	28.6	25.8	35.07	18.26	12.28	8.38	4.05	11.36	13.81	00	00	-	-
NT2RP2005465	8.44	13.41	18.08	10.55	9.11	6.34	5.69	6.77	5.88				
NT2RP2005472	156.47	127.25	179.85	148.2	122.28	139.88	141.88	194.05	159				
NT2RP2005476	18.49	17.23	18.61	18.08	8.51	7.62	10.76	9.58	6.82	00	00	-	-
NT2RP2005498	38.19	48.01	41.98	41.4	60.4	46.27	36.8	25.85	27.88	0	0	-	-
NT2RP2005491	79.87	82.21	87.48	43.7	46.69	33.15	42.26	41.55	39.38	0	00	-	-
NT2RP2005495	8.87	12.67	12.73	8.05	5.23	3.12	3.29	2.22	4.63	0	00	-	-
NT2RP2005496	40.82	33.68	40.13	35.53	16.36	22.89	48.27	41.88	52.81				
NT2RP2005498	14.84	10.07	20.28	8.15	3.71	0.94	3.15	5.06	7.24	0	0	-	-
NT2RP2005501	15.01	11.26	15.06	10.4	6.02	5.78	5.37	7.88	6.8	0	00	-	-
NT2RP2005506	388.46	291.76	402.21	469.83	428.17	208.88	248.04	333.44	241.6				
NT2RP2005509	12.93	15.67	24.29	8.78	9.8	9.01	11.69	14.68	14.73				
NT2RP2005514	15.12	16.77	16.43	9.09	7.1	7.44	5.07	6.13	3.95	00	00	-	-
NT2RP2005520	125.71	88	119.64	107.31	84.69	104.73	102.45	122.23	104.75				
NT2RP2005531	11.61	19.18	16.45	11.91	8.22	5.77	3.77	4.63	4.84	00	00	-	-
NT2RP2005535	6.49	2.54	4.37	5.65	2.65	0.74	1.72	1.38	1.4				
NT2RP2005539	28.78	24.33	27.3	18.42	7.87	16.06	13.78	20.87	16.38	*	0	-	-
NT2RP2005540	7.33	11.84	18.75	8.51	6.39	4.26	3.51	3.93	4.88				
NT2RP2005541	16.33	11.75	18.94	9.77	17.11	5.41	8.79	8.14	4.41	0	0	-	-
NT2RP2005549	14.69	11.15	19.91	13.02	7.16	8.12	7.95	8.57	7	0	0	-	-
NT2RP2005555	11.12	11.81	18.04	9.08	7.83	5.49	4.23	4.44	5.21	0	0	-	-
NT2RP2005555	16.74	21.24	18.63	10.59	10.43	7.67	9.05	5.00	3.87	00	00	-	-
NT2RP2005557	15.1	14.02	12.38	10.08	7.31	4.94	4.63	4.36	4.73	0	00	-	-
NT2RP2005581	28.08	28.29	29.21	28.59	10.92	19.41	31.88	21.24	23.29				
NT2RP2005586	18.13	21.05	17.82	8.84	4.94	6.39	4.8	7.07	6.9	00	00	-	-
NT2RP2005597	6.89	6.87	9.77	7.01	1.89	3.66	2.71	4.02	1.82	0	0	-	-
NT2RP2005600	10.61	7.34	12.5	4.41	8.29	5.16	5.14	5.97	3.47	0	0	-	-
NT2RP2005605	53.68	52.91	55.5	30.27	27.97	29.09	25.09	21.21	26.25	00	00	-	-
NT2RP2005614	9.49	10.07	18.9	6.5	3.69	4.19	2.85	3.64	2.63	0	0	-	-
NT2RP2005620	17.64	10.99	8.05	7.1	2.87	3.18	7.89	3.09	3.37				
NT2RP2005622	16.18	16.55	15.71	6.98	5.85	5.88	5.01	2.43	3.54	00	00	-	-
NT2RP2005632	24.79	21.69	16.89	24.01	11.37	18.72	22.78	10.85	12.69				
NT2RP2005635	41.82	31.3	47.35	47.22	30.84	88.01	30.58	31.82	30.82				
NT2RP2005637	8.32	23.19	12.21	8.57	5.9	4.47	3.6	7.75	5.54				
NT2RP2005640	15.46	12.6	16.31	14.33	9.38	13.4	14.45	17.32	12.67				
NT2RP2005645	23.93	18.08	21.64	10.6	8.02	9.84	11.24	8.09	9.18	00	00	-	-
NT2RP2005651	27.47	30.93	34.21	15.98	19.8	18.12	23.48	20.74	24.88	00	0	-	-
NT2RP2005654	13.13	10.24	14.43	6.38	4.26	2.41	3.58	3.27	5.28	00	00	-	-
NT2RP2005666	12.61	72.48	13.27	7.58	5.9	4.42	7.43	6.32	9.59				
NT2RP2005669	15.23	15	15.44	12.85	8.17	11.55	8.81	7.49	8.17	0	00	-	-
NT2RP2005670	7.59	11.17	18.89	7.08	6.37	8.24	3.98	5.41	8.38	0	0	-	-
NT2RP2005671	6.68	9.38	12.5	6.17	4.27	6.41	3.39	7.42	4.35	0	0	-	-
NT2RP2005675	63.41	58.69	56.21	73.47	47.08	88.08	46.13	47.48	45.27	00	00	-	-
NT2RP2005683	12.5	14.86	22.54	12.03	8.9	11.13	12.61	11.13	11				
NT2RP2005690	7.16	9.78	9.15	7.31	2.68	3.79	6.01	4.76	5.81	0	0	-	-
NT2RP2005694	13.45	13.78	12.27	8.68	4.9	4.48	6.63	6.81	8.01	00	00	-	-
NT2RP2005701	21.28	27.64	24.8	13.54	7.14	7.23	9.98	8.67	7.38	00	00	-	-
NT2RP2005712	5.67	8.5	8.73	7.41	2.31	3.04	4.59	2.88	2.66	0	0	-	-
NT2RP2005719	17.92	12.64	12.02	8.92	5.92	13.08	4.64	11.29	9.06				
NT2RP2005722	24.17	15.94	25.68	22.5	8.47	8.89	8.38	12.95	9.48	0	0	-	-
NT2RP2005723	11.76	19.91	11.37	8.9	4.43	7.12	5.8	6.31	6.59	0	0	-	-
NT2RP2005726	13.16	8.51	18.61	13.17	7.93	10.58	7.85	8.96	10.35				
NT2RP2005729	20.35	17.21	20.34	11.84	8.52	5.78	12.25	8.44	10.44	00	00	-	-
NT2RP2005731	4.25	2.53	4.11	5.17	0	0.69	1.08	1.65	0.86	0	0	-	-
NT2RP2005732	54.72	63.14	65.72	81.89	67.12	83.97	67.17	50.47	67.53	0	+	-	-

82 / 157

2603

2604

NT2RP2005787	61.08	48.51	64.73	67.88	30.52	54.9	80.55	70.19	69.8	*	+
NT2RP2005741	8.17	4.22	11.68	8.13	3	5.42	3.82	4.85	3.85		
NT2RP2005748	8.38	5.27	12.52	8.19	3.71	6.33	3.48	4.61	7.44		
NT2RP2005752	31.76	22.82	38.43	14.99	13.76	18.04	15.51	16.74	18.45	*	-
NT2RP2005753	34.44	30.95	46.77	33.24	28.02	18.11	28.19	26.32	15.55		
NT2RP2005768	8.78	7.82	12.68	8.76	3.85	3.68	4.32	5.6	4.17	*	-
NT2RP2005767	8.89	8.69	15.88	7.42	5.18	5.67	6.13	4.89	5.16		
NT2RP2005773	53.73	61.68	89.19	40.88	32.98	33.21	37.38	30.92	21.68	**	**
NT2RP2005774	13.75	12.5	18.48	16.31	7.06	7.63	18.33	17.56	17.08		
NT2RP2005775	12.13	7.53	22.24	13.79	4.82	6.89	4.04	8.6	6.74		
NT2RP2005781	39.68	23.81	38.05	24.73	22.45	22.65	9.91	16.91	23.08		
NT2RP2005784	30.34	20	41.77	17.54	13.38	14.34	11.13	13.28	15.12		
NT2RP2005789	16.82	16.55	24.71	17.18	13.84	7.39	8.95	14.02	8.78		
NT2RP2005799	11.23	8.75	9.48	10.01	3.28	4.24	3.37	6.03	3.68	**	-
NT2RP2005804	106.7	97.93	105.65	76.82	70.98	67.98	110.4	84.7	108.91	**	-
NT2RP2005812	6.58	9.62	8.38	6.82	4.2	4.89	5.32	2.9	2.58	*	-
NT2RP2005815	2.78	4.88	5.71	6.34	2.94	2.68	4.39	4.52	4.97		
NT2RP2005835	26.87	25.04	37.02	18.31	16.79	18.79	10.34	14.43	39.55	*	-
NT2RP2005841	22.25	27.58	27.52	20.49	14.43	18.92	11.4	11.66	19.61	*	-
NT2RP2005853	18.13	15.89	21.67	12.41	10.18	1.04	9.88	18.61	11.3	*	-
NT2RP2005867	11.03	8.76	10.83	7.42	4.61	1.32	4.21	5.92	5.5	*	-
NT2RP2005859	12.15	11.97	15.23	11.24	7.93	4.88	8.29	8.94	6.43	*	-
NT2RP2005860	0.88	5.77	2.68	4.74	2.31	1.81	1.34	1.77	1.13		
NT2RP2005863	8.01	8.99	7.08	7.46	6.66	5.32	7.67	5.36	4.12		
NT2RP2005868	6.66	10.09	8.76	11.28	4.39	6.75	7.63	5.42	5.14		
NT2RP2005876	28.94	9.38	44.21	26.04	19.27	15.81	14.52	17.33	37.28		
NT2RP2005878	17.4	21.07	28.07	15.88	10.5	10.77	9.96	17.05	21.58		
NT2RP2005883	8.58	9.62	13.3	8.93	8.64	7.61	3.43	11.45	8.28		
NT2RP2005885	109.48	88.27	131.02	108.7	95.74	145.77	77.39	72	85.69		
NT2RP2005887	12.85	18.34	15.15	9.59	5.47	2.19	3.8	6.68	5.07	*	**
NT2RP2005890	11.6	13.16	14.64	6.78	5.32	5.53	4.7	5.77	4.32	**	**
NT2RP2005901	11.86	15.27	14.95	10.18	6.55	8.12	6.42	5	5.8	*	**
NT2RP2005902	4.82	5.49	5.82	9.24	3.89	5.72	4.9	8.33	5.18		
NT2RP2005908	24.41	23.58	59.08	23.37	17.17	15.36	18.41	26.09	27.9		
NT2RP2005927	7.88	6.48	15.44	8.31	4.33	4.48	3.45	5.76	4.29		
NT2RP2005933	14.92	15.67	28.13	12.37	8.74	5	4	6.32	4.28	*	-
NT2RP2005941	10.69	8.58	13.22	7.84	3.9	3.98	2.9	4.82	2.83	*	**
NT2RP2005942	10.35	10.62	11.7	8.52	4	3.48	3.68	4.6	2.9	*	**
NT2RP2005946	8.12	19.6	14.91	5.77	8.58	2.97	4.94	4.86	6.08		
NT2RP2005970	78.5	72.18	87.01	60.77	49.16	40.53	67.44	62.72	62.64	**	*
NT2RP2005980	7.01	5.07	5.9	6.58	2.11	2.14	3.31	3.65	4.51	*	-
NT2RP2005984	12.99	9.14	13.53	7.59	4.09	0.3	2.32	3.71	4.52	*	**
NT2RP2006004	7.54	4.47	13.08	6.53	2.59	2.1	2.72	2.9	3.13		
NT2RP2006013	11.08	6.58	20.3	11.12	5.98	2.47	2.76	11.94	3.04		
NT2RP2006023	79.4	60.83	74.87	51.53	34.84	38.07	64.38	76.12	78.93	*	-
NT2RP2006028	23.13	26.87	22.82	16.89	12.68	6.73	6.52	8.45	15.41	*	**
NT2RP2006038	4.35	4.75	6.83	4.07	1.31	0.83	1.09	0.16	1.4	**	-
NT2RP2006042	16.45	13.74	13.53	9.24	8.66	3.55	5.94	6.22	7.22	*	**
NT2RP2006043	17.81	13.78	15.29	13.7	7.47	8.55	9.41	5.9	6.89	**	-
NT2RP2006052	12.36	8.55	18.5	5.89	3	4.31	2.78	3.7	3.31	*	*
NT2RP2006057	7	8.68	13.05	8.99	3.77	3.67	1.75	3.57	2.95	*	-
NT2RP2006084	31.45	32.66	33.8	26.4	15.66	8.71	8.38	7.35	7.36	*	**
NT2RP2006088	7.85	6.5	16.27	7.3	3.84	3.84	2.88	4.02	3.43		
NT2RP2006089	10.49	8.04	9.64	6.75	2.89	1.65	2.28	1.16	1.94	*	**
NT2RP2006071	12.09	22.31	12.81	10.28	7.27	0.85	3.44	3.17	3.04	*	-
NT2RP2006080	10.61	10.69	22.59	7.82	4.75	0.57	5.24	4.28	5.3		
NT2RP2006092	15.25	7.07	12.08	13.1	8.89	10.62	13.15	8.79	8.66		
NT2RP2006097	52.21	101.2	64.35	48.83	48.31	38.97	25.69	53.43	52.7		
NT2RP2006098	13.67	12.41	21.3	14.85	8.15	8.48	5.13	8.68	3.8	*	-
NT2RP2006099	65.82	60.33	80.23	78.5	49.78	0.24	73.49	73.17	78.2		
NT2RP2006100	14.09	12.17	17.68	9.6	8.02	0.77	4.01	5.07	4.89	**	-
NT2RP2006103	6.26	4.22	7.72	5.83	3.58	2.08	2.74	2.72	2.55	*	-
NT2RP2006106	48	38	56.88	24.88	35.82	35.58	24.01	21.59	18.79	*	-
NT2RP2006127	6.5	4.88	5.96	5.78	1.93	3.45	3.78	1.37	2.06	*	-

83/157

2605

2606

NT2RP2006134	9.57	8.53	12.45	11.76	2.26	6.64	9.28	6.55	5.95
NT2RP2006141	13.0	10.51	20.75	11.91	8.44	5.24	5.17	8.6	15.73
NT2RP2006168	43.19	37.58	47.62	23.75	15.94	0.88	44.65	56.53	70.7
NT2RP2006176	17.92	13.88	20.88	13.87	7.31	8.84	7.42	10.41	8.82
NT2RP2006181	4.75	8.83	6.23	3.92	1.81	1.35	3.97	2.81	2.8
NT2RP2006184	64.52	54.32	69.63	42.14	32.28	42.22	63.72	47.39	43.75
NT2RP2006188	8.19	11.92	9.4	5.19	2.49	2.83	2.18	2.13	1.87
NT2RP2006195	22.53	23.16	21.9	14.52	7.82	10.37	16.26	11.79	10.32
NT2RP2006199	9.01	9.11	6.26	8.72	4.43	4.95	6.19	4.21	5.05
NT2RP2006200	7.28	4.32	9.53	5.05	3.47	3.37	3.59	4.46	2.85
NT2RP2006210	90.78	69.8	59.22	55.44	37.89	25.49	20.22	40.51	30.76
NT2RP2006219	11.28	20.39	17.51	8.1	14.35	6.63	11.34	9.44	19.6
NT2RP2006224	12.75	19.13	21.38	12.72	10.81	3.84	17.87	18.61	1.85
NT2RP2006237	7.47	20.77	9.98	5.08	2.48	0.88	3	2.56	0.6
NT2RP2006238	10.52	14.31	12.03	6.19	1.38	2.03	2.55	2.82	2.68
NT2RP2006258	10.05	20.38	11.9	6.71	3.4	2.58	5.8	4.64	6.64
NT2RP2006261	6.38	9.18	5.31	9.6	2.51	4.78	2.42	4.07	4.58
NT2RP2006280	17.03	14.29	15.32	11	5.59	7.26	3.88	12.08	5.85
NT2RP2006275	48.12	39	46.05	70.83	38.96	80.2	37.5	48.02	38.3
NT2RP2006282	18.4	26.43	24.55	13.47	12.52	6.82	6.66	8.21	6.64
NT2RP2006302	17.18	26.6	32.5	13.26	6.68	8.01	5.28	4.17	5.65
NT2RP2006312	31.59	23.04	41.61	9.42	5.96	4.21	6.1	4.79	7.53
NT2RP2006320	41.28	41.71	38.93	24.89	12.12	15.23	28.98	37.1	32.63
NT2RP2006321	10.41	9.33	12.88	10.28	2.43	5.6	7.6	8.6	8.78
NT2RP2006323	7.58	5.4	4.29	6.82	1.35	3.01	7.41	4.95	4.91
NT2RP2006333	5.03	7.52	4.99	6.18	1.44	5.34	2.1	4.81	3.82
NT2RP2006334	7.94	8.71	7.64	8.5	4.46	8.47	2.3	4.34	3.32
NT2RP2006338	12.7	9.34	18.58	7.32	2.93	4.16	2.16	2.04	3.64
NT2RP2006339	6.41	4.31	4.81	6.37	3.36	3.67	2.31	2.9	2.68
NT2RP2006365	5.78	12.38	3.94	6.81	1.88	2.59	2.74	4.14	2.84
NT2RP2006385	5.27	2.38	4.32	5.39	0.85	1.53	2.13	3.92	2.48
NT2RP2006374	123.01	124.6	151.43	183.82	144.52	148.4	137.08	112.64	168.88
NT2RP2006383	22.69	18.63	23.1	17.79	8.84	13.61	27.87	23.54	25.74
NT2RP2006384	15.6	11.55	17.73	18.97	12.68	13.34	9.59	13.89	11.88
NT2RP2006400	8.82	4.99	12	10.48	4.15	11.24	9.88	11.64	10.9
NT2RP2006411	4.42	4.9	11.08	5.28	2.3	3.07	2.25	3.04	3.21
NT2RP2006429	2.69	1.8	8.38	6.1	1.17	1.68	2.03	2.68	1.48
NT2RP2006435	1.14	1.44	3.88	3.88	0	1.28	1.25	0.25	0.25
NT2RP2006436	18.83	25.85	29.36	23.03	10	15.2	18.88	16.71	17.68
NT2RP2006441	3.24	5.37	8.45	6.78	2.17	3.47	5.6	2.69	1.87
NT2RP2006447	19.2	14.3	18.35	21.17	15.45	16.45	6.09	13.77	11.65
NT2RP2006454	8.91	2.58	13.85	6.26	3.2	3.65	3.95	4.09	2.98
NT2RP2006455	37.87	48.18	39.71	20.34	20.45	22.67	31.23	29.73	37.11
NT2RP2006458	12.66	10.34	20.88	11.22	5.29	5.49	7.54	4.09	3.92
NT2RP2006464	49.31	38.6	70.84	30.42	38.8	32.85	22.89	19.33	15.74
NT2RP2006467	7.41	7.72	12.37	8.13	6.48	4.56	10.4	4.81	8.83
NT2RP2006472	17.27	35.33	39.93	38.59	25.1	29.12	15.84	31.69	29.65
NT2RP2006474	5.31	10.55	8.21	9.45	5.79	7.02	7.91	11.16	8.64
NT2RP2006475	14.67	7.33	18.42	13.4	5.52	5.53	5.81	10.03	7.38
NT2RP2006476	21.17	14.25	23.98	17.93	10.29	15.83	15.59	21.38	17.75
NT2RP2006501	88.84	47.79	67.54	47.85	38.74	38.28	29.87	41.76	52.91
NT2RP2006512	10.73	9.4	12.55	6.68	7.05	5.5	3.87	4.91	5.32
NT2RP2006526	12.09	17.75	22.2	18.92	18.43	13.33	17.31	20.34	13.23
NT2RP2006527	4.41	5.17	7.15	7.48	1.84	1.55	3.51	2.56	2.84
NT2RP2006534	4.37	9.41	7.99	5.05	4.19	2.96	4.58	2.83	2.5
NT2RP2006537	5.9	9.88	11.88	11.28	5.12	7.83	10.34	5.58	8.5
NT2RP2006543	12.11	8.37	13.04	12.23	3.28	11.04	6.8	5.53	9.23
NT2RP2006554	7.01	5.37	13.82	7.47	2.64	2.23	3.56	4.89	9.88
NT2RP2006565	14.34	12.69	22.07	6.94	4.33	4.74	3.36	7.51	9.3
NT2RP2006571	85.34	49.49	69.38	45.6	41.55	34.63	47.88	85.88	43.88
NT2RP2006573	12.88	9.58	16.78	8.82	7.97	5.87	5.7	4.86	5.48
NT2RP2006598	5.82	15.24	10.14	5.39	4.48	3.2	2.52	3.49	4.84
NT2RP2006601	5.82	8.82	7.57	6.45	4.69	2.53	4.15	3.79	3.88
NT2RP2006602	17.28	31.41	14.62	22.94	26.84	18.92	27.08	23.91	17.49

84 / 157

2607

NT2RP3000011	14.9	8	16.29	8.33	4.85	7.79	4.33	10.64	6.28				
NT2RP3000014	108.09	108.54	95.92	88.76	88.18	81.91	47.71	112.2	132.8				
NT2RP3000018	14.35	13.83	24.71	7.65	8.48	5.45	3.86	7.49	4.11	*	*	-	-
NT2RP3000022	7.65	10.24	13.67	9.49	4.92	2.18	2.89	2.32	2.03	*	*	-	-
NT2RP3000024	12.64	12.37	14.37	11.32	5.72	8.81	6.24	6.05	6.62	*	**	-	-
NT2RP3000031	77.24	78.76	145.67	144.42	128.63	86.96	77.23	69.72	83.66				
NT2RP3000034	6.27	6.82	8.74	6.38	3.4	3.43	4.64	3.79	2.88	*	*	-	-
NT2RP3000037	17.58	19.44	20.81	11.72	11.55	13.9	18.45	10.67	14.15	**	*	-	-
NT2RP3000040	27.59	18.23	24.01	13.53	7.79	10.55	2.51	8.1	8.81	*	*	-	-
NT2RP3000041	5.08	4.46	12.76	6.9	3.15	3.65	6.44	3	1.87				
NT2RP3000048	7.83	5.75	16.27	8.29	3.02	2.83	2.71	4.54	5.05				
NT2RP3000047	5.93	4.81	14.06	5.49	2.94	2.87	0.9	3.18	1.81				
NT2RP3000049	6.45	8.2	10.35	6.13	1.6	1.78	2.62	3.99	1.9	*	*	-	-
NT2RP3000050	11.72	13.02	12.95	7.97	7.04	4.74	3.27	4.45	3.83	**	**	-	-
NT2RP3000051	20.55	9.33	16.87	14.09	17.37	4.84	3.99	4.5	4.21	*	*	-	-
NT2RP3000054	8.46	5.67	6.8	6.67	3.62	2.38	1.52	2.23	2.77	**	**	-	-
NT2RP3000055	18.47	13.61	17.33	12.33	8.89	12.37	7.71	11.31	9.35	*	*	-	-
NT2RP3000056	1.87	2.38	8.51	3.35	1.42	1.09	0.81	1.36	0.38				
NT2RP3000059	7.69	7.9	22.29	10.9	7.16	5.13	2.59	4.88	2.31				
NT2RP3000083	1.84	4.14	7.63	4.58	1.09	0.48	0.47	0.97	0				
NT2RP3000088	6.17	8.88	18.21	5.26	4.81	2.29	6.13	4.42	2.89				
NT2RP3000089	9.16	8.31	8.75	6.58	4.58	4.92	4.17	3.76	8.03	**	**	-	-
NT2RP3000072	32.05	40.26	39.16	31.11	26.2	25.34	26.47	26.26	26.37	*	*	-	-
NT2RP3000080	28.2	23.2	22.56	18.48	15.53	13.5	21.79	14.38	13.52	**	*	-	-
NT2RP3000085	67.96	43.89	70.79	53.48	33.71	65.84	52.26	68.62	62.27				
NT2RP3000087	111.9	98.79	117.72	81.88	54.53	62.52	49.82	71.54	66.48	*	**	-	-
NT2RP3000092	9.87	12.43	24.83	8.97	4.99	4.92	2.85	7.38	3.87				
NT2RP3000109	10.57	9.88	12.54	8.31	8.64	5.89	6.65	6.37	5.22	*	**	-	-
NT2RP3000119	8.08	6.85	10.03	6.29	4.73	2.45	4.69	3.89	2.89	*	*	-	-
NT2RP3000125	6.75	8.13	6.71	5.47	2.88	2.09	0.61	2.51	2.19	*	*	-	-
NT2RP3000121	13.6	16.29	13.48	12.62	5.73	6.16	5.37	4.48	4.22	**	**	-	-
NT2RP3000134	18.43	21.97	18.23	9.28	9.9	8.55	6.98	2.89	3.8	**	**	-	-
NT2RP3000137	10.23	18.11	17.39	15.88	4.55	11.71	4.48	6.98	5.44	*	*	-	-
NT2RP3000142	35.53	27.58	35.61	21.77	15.45	18.48	21.05	30.25	26.9	*	*	-	-
NT2RP3000146	26.32	39.38	34.66	19.53	19	15.35	18.54	17.45	14.71	*	*	-	-
NT2RP3000149	32.9	38.83	27.1	14.1	20.28	8.54	10.04	7.73	9.34	*	**	-	-
NT2RP3000183	8.85	9.04	13.21	8.87	3.58	5.95	3.7	3.77	3.35	**	*	-	-
NT2RP3000188	15.39	20.39	19.23	9.48	11.8	5.55	7.63	15.26	9.88	*	*	-	-
NT2RP3000189	15.25	12.33	13.06	7.58	7.27	4.09	8.07	3.27	4.38	**	**	-	-
NT2RP3000171	6.75	12.8	10.82	11.26	6.03	4.95	8.98	2.23	5.04				
NT2RP3000172	40.04	25.12	22.12	25.17	9.97	24.6	26.83	33.18	37.52				
NT2RP3000186	16.1	16.13	19.67	12.39	6.83	8.09	5.65	10.05	10.55	*	*	-	-
NT2RP3000187	50.94	62.27	70	34.47	33.83	20.62	34.62	48.18	55.04	*	*	-	-
NT2RP3000201	14.53	11.88	22.5	7.11	4.59	3.11	5.14	5.78	5.41	*	*	-	-
NT2RP3000204	4.18	4.96	6.69	4.48	1.18	2.3	2.68	1.7	2.27	*	*	-	-
NT2RP3000207	95.69	249.91	234.63	30.21	59.19	29.76	35.62	12.46	31.91	*	*	-	-
NT2RP3000216	7.95	7.31	6.85	6.17	2.69	2.5	2.84	3.29	3.98	*	**	-	-
NT2RP3000220	22.18	29.88	28.92	10.43	6.88	15.01	13.77	5.33	14.12	**	*	-	-
NT2RP3000221	10.31	9.17	11.73	12.25	5.03	10.36	5.52	10.77	8.91				
NT2RP3000232	46.66	42.48	53.25	15.89	8.2	7.9	5.53	11.37	6.25	**	**	-	-
NT2RP3000234	8.3	12.35	2.87	7.16	18.65	2.19	9.24	3.42	2.51				
NT2RP3000234	3.63	3.88	6.88	5.35	3.32	2.04	1.6	2.52	2.2	*	*	-	-
NT2RP3000235	9.45	15.39	13.27	13.07	5.88	8.43	10.72	9.32	10.97				
NT2RP3000239	18.63	25.23	22.23	15.58	11.06	7.77	9.06	9.28	12.71	*	**	-	-
NT2RP3000247	32.05	44.16	41.62	22.89	17.24	20.67	24.3	20.29	19.85	**	*	-	-
NT2RP3000251	44.58	41.2	38.89	25.84	14.62	23.41	43.42	39.09	44.51	**	*	-	-
NT2RP3000252	21.9	16.55	21.84	18.88	16.84	19.17	13.44	23.92	18.04				
NT2RP3000255	18.67	16.02	17.43	18.73	10.58	14.35	9.36	10.73	8.47	**	*	-	-
NT2RP3000262	17.74	15.84	18.49	25.1	18.29	24.75	12.36	12.76	13.38	**	*	-	-
NT2RP3000266	37.8	22.61	54.63	48.81	29.08	37.44	30.75	39.28	44.8				
NT2RP3000267	5.43	5.61	4.39	6.25	5.05	2.74	3.18	2.68	2.59	**	*	-	-
NT2RP3000271	4.82	10.2	10.62	7.08	4.11	3.49	3.88	5.38	2.72				
NT2RP3000276	8.77	11.93	8.85	9.87	5.55	4.79	8.29	7.56	5.48				
NT2RP3000281	32.19	29.94	32.45	16.28	8.76	14.6	16.94	10.41	18.64	**	**	-	-

85 / 157

2608

2609

2610

NT2RP3000292	73.41	53.68	62.68	43.77	42.72	40.88	49.91	45.23	64.29	*	-	-
NT2RP3000299	6.1	3.82	13.07	7.5	3.65	4.11	4.21	4.47	3.84	*	-	-
NT2RP3000304	25.73	24.35	31.49	22.89	17.08	15.95	20.38	9.68	10.08	*	*	-
NT2RP3000310	21.09	16.82	33.28	14.15	13.15	10.27	8.12	6.37	6.54	*	-	-
NT2RP3000312	0.89	6.75	7.52	6.17	1.88	0.95	3.83	3.7	0.45	*	-	-
NT2RP3000320	11.64	11.91	24.03	16.6	10.14	7.39	6.78	13.07	6.3	*	-	-
NT2RP3000322	6.3	10.23	19.74	8.18	5.89	5.4	11.65	9.13	4.6	*	-	-
NT2RP3000324	3.35	4.35	17.78	5.33	3.9	3.91	1.03	4.83	1.94	*	-	-
NT2RP3000328	64.19	41.44	38.63	20.52	18.15	16.6	15.62	30.96	63.08	*	-	-
NT2RP3000329	11.55	11.02	22.68	13.53	7.92	12.17	10.06	11.03	12.62	*	-	-
NT2RP3000330	7.65	5.88	10.92	7.41	2.84	4.65	3.07	4.04	3.9	*	-	-
NT2RP3000333	5.32	5.01	9.94	8.43	4.88	3.17	4.98	3.48	4.27	*	-	-
NT2RP3000341	13.68	14.31	17.67	13.78	9.81	9.9	9.7	9.87	11.82	*	-	-
NT2RP3000344	45.81	52.79	53.7	28.28	29.87	22.15	38.08	31.27	32.79	**	**	-
NT2RP3000345	9.05	22.29	17.79	14.81	9.64	10.42	11.7	6.84	6.21	*	-	-
NT2RP3000348	162.75	78.47	148.51	181.96	103.16	145.4	96.65	108.66	99.32	*	-	-
NT2RP3000350	14.44	9.79	17.94	12.6	7.91	9.89	10.68	16.9	13.81	*	-	-
NT2RP3000359	12.38	12.68	23.75	11.88	10.77	13.58	14.05	17.04	16.61	*	-	-
NT2RP3000381	30.21	20.82	25.69	12.5	10.99	10.9	7.28	12.88	9.35	**	**	-
NT2RP3000360	69.71	53.86	58.74	44.82	38.68	40.33	27.58	41.09	30.61	*	-	-
NT2RP3000378	8.8	12.86	6.1	6.74	3.98	3.58	4.72	5.63	5.68	*	-	-
NT2RP3000384	26.45	19.42	18.46	18.21	9.88	7.79	8.34	6.32	6.93	*	-	-
NT2RP3000389	22.1	25.13	28.16	17.81	18.95	18.45	23.81	18.65	25.81	*	-	-
NT2RP3000393	33.85	14.06	19.19	10.32	8.44	17.78	6.01	7.4	12.32	**	-	-
NT2RP3000395	24.48	28.99	21.1	13.42	18.29	14.47	9.05	14.21	24.87	**	-	-
NT2RP3000397	24.89	21.38	36.91	14.31	14.82	16.14	17.25	23.74	23.2	*	-	-
NT2RP3000398	9.95	10.75	17.31	7.25	6.82	4.6	4.18	4.17	3.99	*	-	-
NT2RP3000403	93.32	64.65	58.89	57.84	43.74	45.83	35.74	40.83	30.88	*	-	-
NT2RP3000418	14.37	22.01	20.2	11.88	12.29	11.51	7.67	14.89	12.11	*	-	-
NT2RP3000424	40.88	56.6	51.72	39.57	32.99	30.24	31.05	28.26	23.73	*	-	-
NT2RP3000427	19.59	24.85	16.69	14.3	18.74	15.64	9.18	9.9	12.52	*	-	-
NT2RP3000431	16.18	14.67	25.82	11.73	18.4	20.33	8.15	16.47	19.08	*	-	-
NT2RP3000433	25.31	54.08	36.8	35.67	31.58	27.85	24.04	39.4	34.85	*	-	-
NT2RP3000436	9.62	10.09	18.49	6.56	4.89	5.29	2.88	5.01	2.75	*	-	-
NT2RP3000439	44.72	47.78	44.4	50.88	60.94	30.43	31.38	44	42.05	*	-	-
NT2RP3000441	10.82	8.95	14.82	10.83	6.19	5.02	6.65	8.63	7.39	*	-	-
NT2RP3000444	17.45	14.75	28.3	14.55	11.35	9.32	12.21	11.85	12.88	*	-	-
NT2RP3000448	6.81	7.47	11.87	6.37	5.01	4.15	4.85	3.6	5.55	*	-	-
NT2RP3000449	17.38	28.19	23.45	8.67	7.34	7.85	3.61	4.2	2.92	**	**	-
NT2RP3000451	31.28	40.71	35.48	10.79	20.02	15.29	13.25	22.78	16.67	**	**	-
NT2RP3000456	8.24	4.2	13.64	8.03	3.84	0	2.84	3.63	2.55	*	-	-
NT2RP3000460	10.33	10.39	17.07	7.27	4.9	3.83	5.19	5.58	4.78	*	-	-
NT2RP3000471	11.88	7.85	28.63	10.17	6.71	3.54	3.88	5.28	6.09	*	-	-
NT2RP3000477	49.73	64.72	58.77	38.09	33.82	22.58	25.2	26.62	28.78	*	-	-
NT2RP3000478	6.95	4.02	8.07	5.51	2.95	1.38	1.83	1.8	1.91	*	-	-
NT2RP3000481	11.03	7.58	11.83	11.71	8.13	6.39	6.85	9.77	6.86	*	-	-
NT2RP3000484	25.46	33.21	34.57	19.58	15.47	13.87	14.97	11.89	12.9	*	-	-
NT2RP3000487	21.14	28.38	33.78	13.82	12.2	14.9	5.33	12.48	12.14	*	-	-
NT2RP3000512	12.44	14.88	14.98	11.55	7.87	5.16	5.05	7.04	6.14	*	-	-
NT2RP3000523	93.94	104.88	80.32	71.84	44.81	35.48	20.12	30.84	18.68	*	-	-
NT2RP3000526	7.84	7.79	12.87	7.19	2.92	3.58	4	5.42	4.09	*	-	-
NT2RP3000527	42.97	38.74	42.72	25	28.15	22.19	19.81	21.07	11.92	**	**	-
NT2RP3000531	29.64	13.82	14.48	12.89	10.75	4.81	8.04	4.79	4.56	*	-	-
NT2RP3000532	8.33	8.65	5.17	4.82	3.38	2.17	2.24	1.62	4.13	*	-	-
NT2RP3000542	21.28	18.32	16.2	8.3	8.11	6.12	3.53	3.83	3.8	**	**	-
NT2RP3000564	11.71	17.87	14.03	9.45	3.02	6.99	4.57	4.9	5.09	*	-	-
NT2RP3000581	21.04	18.88	23.95	14.27	9.57	9.4	4.87	15.18	7.79	*	-	-
NT2RP3000582	66.42	51.6	61.28	47.38	48.18	42.62	32.74	44.25	48.68	*	-	-
NT2RP3000578	19.29	21.22	25	14.6	13.72	9.4	9.07	9.7	10.08	*	-	-
NT2RP3000582	9.4	7.64	13.09	8.29	5.67	6.43	5.12	6.71	6.65	*	-	-
NT2RP3000584	22.71	44.85	35.97	15.47	28.23	10.78	9.11	11.41	6.79	*	-	-
NT2RP3000588	8.23	4.88	7.14	6.37	3.31	2.97	3.81	1.9	2.63	*	-	-
NT2RP3000590	7.21	5.97	5.97	6.82	2.92	1.94	4.82	2.84	1.77	*	-	-
NT2RP3000592	69.63	37.92	80.02	128.75	103.66	125.15	70.02	69.72	60.28	*	+	-

86 / 157

2611

2612

NT2RP3000596	8.71	11.87	13.23	5.92	4.35	3.28	4.65	3.35	4	0	00	-	-
NT2RP3000599	39.3	60.28	67.05	35.1	44.12	24.38	35.82	38.21	35.98				
NT2RP3000603	69.47	91.4	87.14	91.6	88.66	93.49	64.94	61.32	72.5				
NT2RP3000605	22.45	27.45	25.73	13.14	8.89	13.48	17.63	11.14	9.78	00	0	-	-
NT2RP3000607	10.48	11.53	16.41	10.77	7.31	9.58	5.29	9.79	8.32				
NT2RP3000616	11.88	13.13	17.29	7.98	6.83	4.22	6.69	4.25	4.27	0	00	-	-
NT2RP3000621	218.49	188.17	248.48	222.59	168.47	221.1	202.08	168.53	188.24				
NT2RP3000622	11.89	8.26	8.27	9.18	8.89	5.28	8.73	8.99	3.57				
NT2RP3000624	27.07	18.6	22.66	18.72	12.03	13.8	17.33	23.33	18.89	*		-	-
NT2RP3000628	27.69	34.41	79.45	22.73	20.51	22.18	16.98	19.75	18.56				
NT2RP3000631	6.03	12.91	11.07	11.79	9.81	5.14	4.52	6.12	7.95				
NT2RP3000632	16.5	21.72	26.01	18.28	13.03	18.88	15.73	14.14	13.67				
NT2RP3000638	31.94	42.93	38.07	22.78	17	20.23	24.56	32.44	30.42	00		-	-
NT2RP3000644	26.66	26.83	24.84	18.68	14.97	15.78	28.13	27.03	30.11	00		-	-
NT2RP3000645	12.37	13.85	21.98	14.87	10.1	8.68	7	6.44	7.2				
NT2RP3000652	20.17	23.26	25.45	12.87	9.39	13.28	9.11	9.19	16.37	00	0	-	-
NT2RP3000658	6.96	4.89	8.62	8.05	2.7	4.41	2.18	3.59	3.85				
NT2RP3000660	21.84	12.21	11.63	11.38	9.42	9.58	9.02	8.73	8.05				
NT2RP3000661	15.39	18.23	27.58	11.62	4.78	9.25	6.28	7.03	6.07	0	0	-	-
NT2RP3000665	34.84	43.07	59.18	27.69	16.88	21.15	18.2	20.9	14.32	0	0	-	-
NT2RP3000676	11.63	4.79	8.95	8.8	4.94	2.4	4.94	2.78	3.68				
NT2RP3000677	12.19	14.6	7.18	13.97	4.51	9.65	9.82	11.58	12.08	0	00	-	-
NT2RP3000681	7.72	7.98	7.04	8.12	2.54	4.33	4.62	3.23	3.28				
NT2RP3000682	16.43	8.63	13.69	14.68	12.94	13.49	5.83	8.27	7.59				
NT2RP3000685	8.41	6.48	7.19	10.24	3.04	4.6	5.55	3.8	3.45				
NT2RP3000686	11.62	7.65	10.23	15.03	9.13	13.25	14.85	17.96	14.92	0		+	
NT2RP3000688	6.94	6.35	7.8	8.41	2.13	5.83	3.39	7.61	6.51				
NT2RP3000708	9.74	10.33	12.77	11.25	8.14	10.39	9.73	10.17	12.88				
NT2RP3000718	9.88	16.24	14.31	8.23	2.38	6.7	5.89	3.68	4.39	0	0	-	-
NT2RP3000721	8.23	3.7	5.33	7.95	1.84	5.33	8.54	6.47	6.13				
NT2RP3000728	4.68	4.59	4.66	7.18	1.44	3.15	5.16	3.82	3.57				
NT2RP3000730	14.89	15.22	17.78	18.78	11.05	11.89	16.08	10.04	9.6				
NT2RP3000733	62.48	50.63	61.88	55.7	39.3	45.85	39.59	39.01	40.88				
NT2RP3000735	4.91	3.22	9.89	8.52	1.99	3.91	2.03	3.65	2.09				
NT2RP3000736	6.38	3.45	10.88	8.89	2.09	4.11	3.75	3.03	2.58				
NT2RP3000739	40.42	44.03	40.83	24.77	27.39	34.23	25.37	20.3	32.14	*	0	-	-
NT2RP3000742	4.84	6.47	8.58	7.03	4.16	2.9	3.28	5.74	3.55				
NT2RP3000753	8.3	6.8	12.68	10.6	4.35	5.47	5.99	5.75	7.45				
NT2RP3000759	38.84	70.83	56.01	36.02	18.41	27.88	8.09	6.25	5.44	00		-	-
NT2RP3000789	45.64	27.39	34.07	33.12	27.39	34.64	20.51	27.87	27.21				
NT2RP3000815	22.33	13.62	22.75	12.02	10.68	14.79	18.74	20.82	19.98				
NT2RP3000818	30.57	21.11	33.63	24.24	17.69	21.68	13.73	17.32	15.5	0		-	-
NT2RP3000820	36.87	28.12	43.82	32.81	27.98	25.05	21.15	25.93	23.2				
NT2RP3000821	3.61	4.2	7.15	6.21	2.3	3.01	3.82	2.8	1.62				
NT2RP3000825	18.22	23.18	33.6	11.87	17.67	13.25	9.13	8.99	18.22				
NT2RP3000826	57.22	56.7	80.61	55.4	48.67	48.17	57.78	64.97	54.17				
NT2RP3000838	8.1	3.53	3.85	7.53	2.33	2.42	3.89	4.81	2.68				
NT2RP3000838	14.68	14.22	18.83	20.82	12.45	10.78	11.08	17.41	54.13				
NT2RP3000839	45.25	39.9	44.36	35.78	27.32	20.79	19.28	33.14	25.16	0	0	-	-
NT2RP3000841	12.21	10.37	21.1	12.42	10.04	9.18	8.34	11.21	7.12				
NT2RP3000845	8.69	31.26	15.5	9.69	12.93	10.25	6.37	12.21	8.92				
NT2RP3000847	12.91	10.6	18.85	11.08	8.9	9.57	6.7	5.47	6.54	*		-	-
NT2RP3000848	11.47	16.2	11.34	8.1	7	4.83	3.86	7.4	3.83	0	0	-	-
NT2RP3000850	21.09	31.1	29.31	19.76	12.79	16.25	10.7	10.54	5.19	0	00	-	-
NT2RP3000852	2.73	3.9	3.17	7.61	3.17	2.39	4.15	2.97	3.68				
NT2RP3000859	32.3	21.43	36.02	18.9	21.77	13.31	18.61	18.23	32.52				
NT2RP3000861	15.66	12.49	16.59	8.02	5.38	4.49	4.53	5.88	27.25	00		-	-
NT2RP3000862	12.15	10.89	17.99	10.38	8.14	8.17	5.59	8.92	5.21				
NT2RP3000865	9.02	8.17	10.88	7.46	4.29	4.01	2.73	4.35	3.2	0		-	-
NT2RP3000866	43.48	38.1	57.47	39.52	40.18	44.87	37.48	37.57	27.68				
NT2RP3000868	10.65	23.22	17.15	7.33	6.65	4.58	6.64	9.1	10.81	0		-	-
NT2RP3000869	5.78	11.8	14.52	9.41	10.88	7.13	8.87	5.06	7.12				
NT2RP3000871	184.48	180.18	141.72	262.18	170.88	170.92	247.35	193.7	214.48				
NT2RP3000875	14.37	14.48	23.84	13.76	12.05	11.47	7.45	6.27	6.52	0		-	-

87 / 157

2613

2614

WT2RP3000895	2.84	2.51	12.27	4.16	1.17	2.03	1.28	3.4	1.59										
WT2RP3000900	10.4	8.3	14.31	16.92	3.45	5.38	4.15	13.51	4.83										
WT2RP3000901	14.3	18.5	22.78	13.03	13.47	11.77	11.97	9.53	14.53										
WT2RP3000903	5.01	6.85	6.69	5.82	1.64	3.35	1.28	1.82	1.35										
WT2RP3000904	108.6	100.47	123.83	118.07	108.3	100.81	97.74	77.69	108.28										
WT2RP3000907	84.64	80.19	89.68	84.77	80.01	92.59	125.26	82.21	80.83										
WT2RP3000913	19.19	19.49	20.62	24.35	8.67	18.44	17.39	14.75	19.21										
WT2RP3000917	18.28	18.3	30.01	16.98	15.22	12.11	11.68	16.53	14.68										
WT2RP3000919	9.33	7.85	14.92	9.52	7.41	6.8	4.48	8.2	5.92										
WT2RP3000921	112.89	136.59	133.65	118.03	103.41	74.11	25.71	35.79	29.81										
WT2RP3000942	57.7	68.21	88.84	75.24	51.25	30.13	38.31	52.76	48.83										
WT2RP3000968	50.83	81.86	37.11	29.89	24.82	14.88	9.88	11.81	12.71										
WT2RP3000974	8.98	8.74	9.35	9.04	12.66	5.51	4.39	4.92	2.99										
WT2RP3000980	52.51	65.55	64.13	24.76	40.04	30.49	19.39	8.08	15.43										
WT2RP3000984	21.27	24.19	17.57	32.72	12.14	21.58	18.47	14.77	17.87										
WT2RP3000994	13.76	10.35	18.12	10.02	5.77	7.25	4.5	8.07	7.2										
WT2RP3001001	26.43	28.79	25.4	17.28	14.54	13.11	11.19	38.41	33.45										
WT2RP3001004	4.86	4.98	10.62	7.73	3.07	4.03	3.13	4.4	2.41										
WT2RP3001007	7.85	8.86	18.59	12.13	9.28	3.44	5.05	8.53	5.77										
WT2RP3001012	29.74	22.53	38.08	15.24	16.48	13.42	12.28	11.59	12.53										
WT2RP3001042	8.82	8.59	12.15	8.9	5.69	3.5	3.15	4.09	3.85										
WT2RP3001044	9.07	11.65	10.99	8.25	8.28	5.69	7.83	7.51	5.88										
WT2RP3001048	6.16	7.8	4.48	10.09	2.76	3.93	4.91	5.32	6.21										
WT2RP3001050	13.37	10.61	15.08	10.7	4.83	9.6	6.34	11.28	8.4										
WT2RP3001055	18.05	10.48	21.89	13.3	7.76	7.17	10.39	12.83	10.28										
WT2RP3001087	402.89	324.38	250.29	272.31	285.99	209.38	215.17	328.65	144.83										
WT2RP3001091	7.38	5.55	10.57	7.58	3.5	2.09	3.87	3.84	2.18										
WT2RP3001089	5.39	4.71	9.8	4.84	1.4	2.62	2.39	2.57	3.49										
WT2RP3001074	12.1	8.82	12.25	8.96	3.61	6.85	9.26	8.39	6.94										
WT2RP3001078	13.01	11.28	14.67	14.05	8.24	8.9	10.46	7.65	9.48										
WT2RP3001081	8.54	6.08	6.98	10.22	3.89	7.03	6.21	3.36	4.4										
WT2RP3001084	37.48	37.33	40.93	15.51	10.29	21.45	23.22	30.25	29.17										
WT2RP3001085	7.78	24.69	13.38	9.04	3.7	5.9	3.38	7.52	4.23										
WT2RP3001086	10.65	10.08	12.53	7.44	5.06	4.18	3.27	5.68	3.18										
WT2RP3001087	16.83	38.99	30.48	15.02	9.07	10.26	12.7	8.94	6.97										
WT2RP3001107	28.61	24.48	44.48	22.05	25.18	18.26	10.04	8.83	11.02										
WT2RP3001109	17.96	17.48	18.98	10.7	8.3	6.68	7.73	6.5	3.91										
WT2RP3001111	10.67	15.58	14.57	9.33	6.41	6.38	8.41	3.97	4.53										
WT2RP3001112	31.69	26.22	26.11	22.8	9.31	16.42	16.88	8.38	14.31										
WT2RP3001113	14.44	9.43	17.81	10.89	5.21	10.46	6.16	6.79	9.04										
WT2RP3001115	254.58	276.58	258.3	135.39	153.31	123.7	118.47	299.67	339.02										
WT2RP3001118	5.38	8	8.83	6.23	4.02	2.69	4.29	3.33	3.99										
WT2RP3001119	18.64	22.88	19.35	7.26	6.2	5.92	8.84	7.37	9.08										
WT2RP3001120	6.48	9.54	10.55	5.21	2.08	2.39	3.48	2.35	2.85										
WT2RP3001126	20.13	19.18	15.24	8.77	8.2	4.55	4.48	2.87	5.52										
WT2RP3001127	348.14	574.68	264.45	161.21	220.83	188.52	583.89	289.72	342.5										
WT2RP3001133	93.42	110.08	112.43	155.23	89.98	121.37	108.06	85.85	85.22										
WT2RP3001140	88.84	96.81	111.2	55.83	48.1	51.01	55.81	103.28	84.66										
WT2RP3001147	4.72	7.64	8.71	8.37	2.09	3.65	2.74	3.68	2.93										
WT2RP3001150	11.47	12.3	12.64	9.68	10.24	3.21	4.96	7.94	6.34										
WT2RP3001152	38.54	72.08	61.62	86.57	55.9	58.11	48.19	41.5	53.58										
WT2RP3001155	19.7	23.76	17.71	11.79	7.15	8.62	14.02	11.36	10.16										
WT2RP3001156	19.48	20.53	21.5	20.77	20.49	12.06	20.09	17.5	18.2										
WT2RP3001159	18.65	19.29	22.76	15.71	8.43	11.51	18.86	20.4	20.19										
WT2RP3001170	16.8	14.88	12.69	11.85	4.5	6.68	9.82	8.85	4.53										
WT2RP3001175	17.98	17.32	14.24	8.81	7.88	12.88	9.17	14.49	11.22										
WT2RP3001195	197.73	229.27	180.42	289.15	260.22	386.11	194.82	191.1	184.18										
WT2RP3001208	74.83	87.88	81.84	88.53	60.41	80.53	81.32	84.97	83.11										
WT2RP3001214	17.39	13.69	27.12	17.15	13.89	18.51	7.87	14.21	13.2										
WT2RP3001215	18.74	16.38	19.13	19.54	12.84	10.16	26.01	26.17	21.52										
WT2RP3001221	6.31	7.39	8.65	9.62	5.18	6.37	8.56	6.05	7.92										
WT2RP3001226	22.75	26.78	31.18	8.49	9.07	6.78	11.49	9.21	9.9										
WT2RP3001230	8.51	8.58	3.89	9.46	3.93	5.31	7.3	8.36	5.9										
WT2RP3001232	6.16	3.81	12.04	7.09	3.34	3.66	1.97	4.41	3										

88 / 157

2615

2616

WT2RP3001236	30.11	10.17	30.67	17.48	11.33	17.06	7.88	12.61	12.49	*	-
WT2RP3001239	12.50	7.93	16.69	7.8	5.43	5.82	7.31	5.36	3.8	*	-
WT2RP3001240	16.64	15.42	32.60	20.34	13.67	11.75	7.09	11.44	8.12	*	-
WT2RP3001245	8.04	12.54	22.57	10.74	8.24	10.53	7.27	7.75	6.31	*	-
WT2RP3001253	5.41	9.93	13.75	5.38	2.74	1.6	4.68	3.73	2.31	*	-
WT2RP3001259	278.9	310.98	127.25	256.7	262.64	274.28	334.89	289.89	242.91	*	-
WT2RP3001260	18.94	14.44	22.33	34.17	17.19	23.22	21.98	25.77	20.63	*	-
WT2RP3001264	67.04	56.89	77.24	82.42	71.22	82.52	48.6	44.18	61.74	*	-
WT2RP3001268	54.13	43.25	63.3	45.53	25.93	35.18	39.03	48.28	54.07	*	-
WT2RP3001271	7.61	5.79	12.88	6.16	2.76	3.69	3.87	7.7	5.38	*	-
WT2RP3001272	28.18	22.6	31.48	18.93	15.25	15.77	18.56	19.38	20.08	*	-
WT2RP3001274	149.91	121	184.35	135.41	115.24	154.69	133.05	136.37	139.51	*	-
WT2RP3001275	7.03	7.19	20.08	7.2	8.45	4.21	4.17	4.56	3.94	*	-
WT2RP3001280	14.64	18.72	19.46	9.93	11.17	5.67	7.6	8.91	6.48	*	-
WT2RP3001281	0.38	2.63	0.7	4.71	1.73	1.17	1.88	0.77	0.6	*	-
WT2RP3001288	43.86	21.34	42.34	25.62	18.43	26.29	27.01	32.19	37.1	*	-
WT2RP3001297	77.1	64.75	81.83	35.65	27.23	33.6	12.09	28.24	29.77	**	-
WT2RP3001300	9.48	9.08	13.61	7.61	5.56	4.68	4.6	5.82	8.28	*	-
WT2RP3001301	5.4	3.52	7.37	3.86	1.62	1.28	1.65	3.26	1.29	*	-
WT2RP3001307	13.22	10.36	15.55	10.66	10.08	6.04	5.09	8.17	7.35	*	-
WT2RP3001310	9.88	12.87	18.58	11.65	7.95	6.71	4.27	4.42	3.89	*	-
WT2RP3001318	7.4	8.54	9.21	7.25	6.79	6.73	5.34	5.32	3.43	*	-
WT2RP3001322	11.09	11.88	14.04	15.94	12.19	11.47	14.68	6.17	7.73	*	-
WT2RP3001325	1291.7	1025	1099	288.93	377.86	405.84	75.05	206.06	218.63	**	-
WT2RP3001338	182.73	132.64	198.98	171.74	181.83	241.04	134.27	181.16	143.19	*	-
WT2RP3001339	18.04	11.7	17.8	11.35	7.84	6.68	2.83	5.64	4.58	*	-
WT2RP3001340	35.84	31.42	47.79	30.03	31.99	45.28	31.64	27.19	27.6	*	-
WT2RP3001341	37.44	66.22	47.17	33.31	24.98	24.34	25.96	38.56	34.37	*	-
WT2RP3001354	42.85	32.23	43.99	23.31	23.66	19.15	13.01	13.77	11.4	*	-
WT2RP3001355	2.26	7.76	10.15	6.3	3.13	3.72	4.68	3.96	2.91	*	-
WT2RP3001356	43	54.12	51.32	82.98	37.84	46.81	81.45	50.61	59.12	*	-
WT2RP3001359	34.72	29.05	42.17	41.55	33.32	37.07	21.88	23.34	24.66	*	-
WT2RP3001364	19.28	26.48	29.33	17.88	11.52	13.89	13.78	18.18	15.95	*	-
WT2RP3001373	18.79	12.24	26.6	10.58	10.1	9.34	9.17	17.19	12.49	*	-
WT2RP3001374	95.59	73.12	108.77	138.19	114.68	107.19	68.33	56.48	69.32	*	-
WT2RP3001383	12	18.08	15.5	12.1	5.75	6.51	13.27	8.62	8.62	*	-
WT2RP3001384	22.91	17.21	30.7	33.57	22.7	20.56	19.13	14.74	15.12	*	-
WT2RP3001388	9.38	9.17	7.78	9.06	5.69	4.27	7.73	5.71	5.18	*	-
WT2RP3001392	4.64	4.2	2.65	5.43	2.03	2.78	2.37	2.94	3.1	*	-
WT2RP3001395	47.27	37.4	60.04	23.62	14.23	34.71	12.93	30.54	38.42	*	-
WT2RP3001398	13.09	9.85	15.48	8.92	6.22	7.1	4.4	8.04	8.53	*	-
WT2RP3001399	17.01	14.49	24.37	18.06	7.16	8.8	8.07	7.51	6.82	*	-
WT2RP3001402	30.22	31.51	33.24	60.13	23.85	9.6	15.21	24.77	23.82	*	-
WT2RP3001407	72.56	60.88	88.74	88.01	74.06	85.12	43.81	83.39	81.28	*	-
WT2RP3001416	18.68	22.22	16.43	16.79	17.16	5.73	3.31	4.98	5	**	-
WT2RP3001420	5.68	3.11	4.79	4.91	2.62	1.74	1.34	1.67	2.52	*	-
WT2RP3001425	11.74	10.47	11.8	11.91	6.37	9.11	8.4	4.94	6.04	*	-
WT2RP3001426	12.17	11.79	16.88	9.1	9.84	6.43	5.51	9.62	7.61	*	-
WT2RP3001427	18.61	11	21.59	12.3	6.43	8.87	17.63	22.78	18.89	*	-
WT2RP3001428	7.7	10.58	20.18	8.58	3.38	6.04	5.05	8.84	8.9	*	-
WT2RP3001429	4.55	8.04	12.09	5.05	1.79	1.93	2.26	1.46	1.89	*	-
WT2RP3001432	29.32	18.99	30.98	16.39	14.24	12.25	16.49	28.23	11.69	*	-
WT2RP3001439	31.43	37.32	28.95	25.93	25.44	19.21	14.96	11	10.21	**	-
WT2RP3001441	34.77	41.28	46.38	53.64	42.72	50.38	30.69	23.25	24.74	*	-
WT2RP3001446	4.24	2.49	4.94	4.63	3.01	2.2	2.82	2.01	1.51	*	-
WT2RP3001447	22.15	21.24	25.11	13.3	8.65	15.94	13.67	37.08	14.99	*	-
WT2RP3001449	139.79	126.76	203.02	199.4	188.47	178.45	131.32	104.74	139.38	*	-
WT2RP3001453	31.6	35.5	49.43	34.98	29.3	38.04	28.49	31.86	35.69	**	-
WT2RP3001457	13.38	17.23	17.13	11.75	10.12	8.15	5.88	6.75	8.92	*	-
WT2RP3001459	11.29	13.84	13.09	7	3.78	5.3	5.53	4.44	5.41	**	-
WT2RP3001463	8.65	4.84	3.91	4.67	1.94	1.07	1.96	2.08	0.69	*	-
WT2RP3001464	5.28	4.9	5.4	5.77	2.61	1.65	2.18	1.94	1.08	**	-
WT2RP3001472	5.37	5.28	7.05	12.24	7.01	3.69	1.12	2.87	2.9	*	-
WT2RP3001475	230.39	301.8	150.57	92.73	106.13	96.77	29.64	88.86	93.78	*	-

89 / 157

2617

2618

WT2RP3001470	6. 12	3. 77	10. 27	6. 45	4. 18	2. 2	2. 6	4. 58	5. 3
WT2RP3001480	15. 72	6. 25	23. 45	15. 28	5. 31	6. 47	7. 24	7. 84	8. 84
WT2RP3001492	14	14. 91	21. 99	13. 78	7. 53	8. 81	14. 28	13. 21	12. 48
WT2RP3001495	18. 58	21. 11	19. 03	12. 08	9. 76	9. 93	15. 89	18. 33	21. 35
WT2RP3001497	38. 57	27. 99	40. 31	47. 7	28. 76	38. 87	23. 69	21. 43	24. 3
WT2RP3001501	16. 78	28. 67	29. 3	12. 62	8. 54	7. 31	11. 70	2. 93	9. 87
WT2RP3001527	5. 64	5. 58	5. 58	10. 27	7. 34	2. 85	3. 31	4. 71	3
WT2RP3001529	21. 9	27. 01	22. 29	20. 64	12. 59	18. 78	7. 77	18. 39	18. 96
WT2RP3001538	80. 08	67. 26	78. 24	69. 51	76. 4	114. 47	59. 99	77. 08	72. 65
WT2RP3001539	4. 24	4. 99	8. 09	6. 1	3. 34	4. 14	3. 58	5. 72	3. 56
WT2RP3001542	17. 53	25. 89	26. 44	14. 74	14. 15	8. 68	14. 26	12. 41	17. 97
WT2RP3001549	8. 14	6. 2	5. 77	5. 78	1. 73	1. 92	2. 67	2. 05	1. 89
WT2RP3001554	28. 4	40. 68	42. 76	10. 47	10. 16	7. 96	15. 82	8. 06	13. 22
WT2RP3001560	79. 18	91. 03	88. 06	84. 47	48. 32	78. 97	115. 31	93. 82	84
WT2RP3001561	3. 47	3. 93	3. 28	6. 28	2. 06	3. 29	2. 23	2. 16	2. 23
WT2RP3001564	34. 74	32. 87	34. 89	17. 25	16. 65	19. 41	10. 11	30. 58	15. 49
WT2RP3001568	23. 08	14. 44	30. 67	16. 82	14. 73	6. 11	7. 97	12. 05	7. 85
WT2RP3001575	34. 43	18. 91	39. 6	19. 69	12. 66	21. 53	16. 47	22. 21	25. 65
WT2RP3001580	17. 53	21. 3	27. 37	11. 9	7. 8	11. 26	7. 5	7. 41	8. 67
WT2RP3001587	14. 32	16. 9	20. 18	11. 06	6. 8	8. 98	19. 28	16. 35	20. 31
WT2RP3001589	6. 31	5. 44	5. 39	5. 29	2. 4	2. 48	2. 34	4. 87	3. 41
WT2RP3001592	3. 73	61. 22	57. 18	30. 41	23. 17	28. 18	46. 3	38. 68	32. 65
WT2RP3001607	11. 49	11. 68	9. 82	7. 34	3. 49	8. 45	5. 15	2. 53	3. 96
WT2RP3001608	10. 61	24. 16	28. 88	15. 14	19	18. 45	6. 66	10. 92	10. 91
WT2RP3001813	55. 34	38. 31	54. 71	81. 86	48. 17	86. 77	51. 11	48. 11	44. 78
WT2RP3001819	162. 64	181. 38	185. 09	152. 03	118. 43	125. 65	144. 48	157. 98	158. 08
WT2RP3001821	29. 37	24. 25	25. 20	33. 83	16. 83	30. 34	32. 22	27. 66	24. 1
WT2RP3001829	28	13. 33	26. 18	16. 26	12. 69	18. 65	19. 79	23. 48	12. 56
WT2RP3001930	3. 51	2. 38	4. 56	4. 63	1. 01	2. 1	3. 79	2. 64	3. 87
WT2RP3001931	64. 16	51. 12	67. 22	112. 91	65. 5	60. 67	59. 38	50. 82	71. 84
WT2RP3001934	3. 71	4. 65	2. 84	7. 65	2. 25	2. 09	4. 82	2. 51	1. 18
WT2RP3001942	20. 24	22. 78	34. 25	25. 21	18. 32	21. 48	34. 78	35. 92	40. 14
WT2RP3001946	10. 72	13. 31	23. 84	17. 34	7. 48	8. 49	11. 13	12. 12	12. 05
WT2RP3001950	4. 68	6. 33	12. 48	7. 77	6. 89	4. 37	3. 47	3. 76	3. 82
WT2RP3001967	17. 36	19. 67	39. 78	21. 64	7. 89	10. 07	11. 81	14. 81	13. 16
WT2RP3001971	13. 32	10. 22	13. 88	10. 05	5. 86	9. 27	7. 9	8. 32	3. 97
WT2RP3001976	68. 18	72. 22	97. 84	28. 72	18. 18	21. 65	10. 7	4. 04	14. 85
WT2RP3001978	11. 84	10. 56	12. 76	8. 88	5. 99	6. 98	7. 09	2. 31	6. 88
WT2RP3001979	358. 4	207. 45	268. 13	288. 3	249. 2	237. 9	189. 86	337. 31	274. 18
WT2RP3001980	19. 45	8. 41	17. 68	13. 47	6. 77	7. 99	3. 71	6. 64	7. 99
WT2RP3001982	116. 37	82. 43	118. 94	71. 89	86. 26	91. 02	56. 88	114. 15	137. 5
WT2RP3001985	4. 91	3. 52	11. 94	5. 18	2. 78	2. 73	5	2. 99	2. 91
WT2RP3001988	4. 75	1. 41	4. 66	3. 33	0. 27	0. 64	0. 89	2. 07	0
WT2RP3001990	8. 33	16. 1	11. 83	11. 07	4. 69	5. 02	5. 74	5. 13	6. 32
WT2RP3001992	2. 22	8. 17	10. 5	5. 55	2. 66	2. 65	2. 84	2. 33	2. 39
WT2RP3001998	7. 34	11. 29	11. 31	6. 93	5. 94	11. 43	6. 97	3. 31	6. 44
WT2RP3001999	780. 45	572. 31	725. 4	616. 13	658. 06	0. 94	533. 83	1241. 6	1289. 4
WT2RP3001708	348. 43	237. 64	257. 74	514. 57	355. 27	1. 82	148. 2	628. 13	510. 33
WT2RP3001712	32. 9	8. 27	35. 65	18. 32	14. 72	16. 28	22. 44	32. 16	22. 67
WT2RP3001716	13. 86	9. 55	15. 81	8. 27	11. 28	8. 27	5. 55	15. 45	9. 22
WT2RP3001724	28. 26	21. 7	18. 16	32. 47	31. 93	26. 73	18. 65	28. 64	18. 15
WT2RP3001727	27. 92	15. 82	33. 03	16. 01	12. 47	12. 24	8. 97	8. 14	8. 89
WT2RP3001729	2. 64	5. 64	6. 12	6. 09	2. 61	2. 66	2. 5	2. 88	1. 85
WT2RP3001730	421. 81	624. 98	490. 81	328. 42	274. 18	316. 08	179. 32	117. 74	101. 94
WT2RP3001733	1. 54	1. 62	4. 9	4. 53	0	1. 49	0. 54	0. 38	1. 87
WT2RP3001737	7. 85	25. 52	17. 42	10. 78	4. 14	4. 59	3. 35	7. 04	17. 38
WT2RP3001738	28. 07	24. 45	41. 38	18. 75	13. 82	15. 18	12. 56	15. 63	32. 84
WT2RP3001742	2. 19	2. 42	9. 18	4. 78	1. 8	1. 62	1. 08	2. 07	0. 9
WT2RP3001751	1. 1	11. 43	14. 41	9. 77	6. 51	6. 96	7. 73	6. 52	8. 12
WT2RP3001752	1. 19	3. 87	3. 32	5. 63	0. 3	1. 67	1. 29	0. 92	0. 82
WT2RP3001752	2. 48	4. 8	2. 87	3. 47	0. 21	1. 52	1. 75	2. 38	0. 9
WT2RP3001753	3. 95	5. 83	5. 77	7. 82	4. 31	2. 88	4. 73	5. 28	2. 51
WT2RP3001754	3. 14	1. 28	9. 55	3. 89	0. 47	2. 48	0. 89	1. 82	0. 66
WT2RP3001766	4. 67	4. 58	13. 42	1. 12	2. 77	3. 58	2. 82	9. 4	3. 58

2619

2620

WT2RP3001764	1.84	4.87	12.85	4.07	1.87	1.79	1.19	3.02	2.08
WT2RP3001771	10.53	6.71	15.52	12	8.69	4.55	2.42	4.06	4.07
WT2RP3001777	40.85	47.70	33.95	47.7	28.57	21.83	29.91	18.82	40.99
WT2RP3001782	18.47	18.59	14.68	13.14	8.87	10.49	5.89	13.23	8.58
WT2RP3001787	51.98	43.73	63.18	42.18	33.84	33.35	80.48	61.45	65.58
WT2RP3001799	7.6	9.23	9.91	6.52	4.07	3.35	7.42	5.61	3.54
WT2RP3001819	9.21	8.05	11.94	7.62	3.17	5.23	2.67	7.27	0.12
WT2RP3001829	0.79	0.98	6.7	3.8	0.17	1.25	3.23	1.13	0.11
WT2RP3001836	16.84	18.28	30.32	14.13	7.32	15.63	6.9	7.75	6.48
WT2RP3001839	6.81	9.19	13.78	8.8	4	4.34	2.3	5.79	3.64
WT2RP3001844	5.94	4.03	11.89	5.39	1.37	2.09	2.04	1.24	0.13
WT2RP3001848	53.13	36.02	38.27	40.15	39.28	28.02	48.85	51.68	51.6
WT2RP3001854	54.05	46.79	63.64	47.02	38.11	28.76	72.37	61.23	74.82
WT2RP3001855	8.77	9.33	11.43	8.53	8.22	8.62	9.1	7.34	5.5
WT2RP3001857	2.78	1.95	7.57	5.8	0.8	2.83	1.63	2.14	0.86
WT2RP3001858	1.48	1.77	7.65	6.01	1.18	1.85	0	1.41	1.37
WT2RP3001861	4.68	3.78	12.83	6.09	2.09	1.64	1.75	2.42	0.53
WT2RP3001866	4.29	4.19	8.58	7.28	2.38	3.86	2.27	2.92	0.73
WT2RP3001871	4.11	2.37	11.14	4.75	0.71	0.4	0.62	1.38	0.06
WT2RP3001874	10.44	6.88	11.3	5.07	2.46	1.93	2.08	3.65	5.3
WT2RP3001878	8.84	8.82	7.26	6.09	2.26	2.81	4.01	2.14	3.48
WT2RP3001885	6.75	5.91	9.28	7.33	5.82	5.28	4.29	3.88	3.88
WT2RP3001886	16.8	9.61	18.78	10.19	6.77	12.23	2.88	7.21	8.15
WT2RP3001893	10.12	6.81	19.59	9.29	3.83	4.48	2.35	4.38	3.15
WT2RP3001899	4.48	3.48	9.85	6.74	4	4.77	3.33	4.48	3.02
WT2RP3001901	2.97	2.49	8.61	4.80	2.11	2.05	1.6	0.22	0.74
WT2RP3001915	21.98	18.4	24.23	18.83	15.48	12.75	6.48	9.55	5.88
WT2RP3001926	42.5	70.22	59.74	48.58	37.18	35.96	45.08	32.42	25.88
WT2RP3001929	7.68	7.59	9	5.33	1.89	5.78	3.26	1.47	1.63
WT2RP3001931	7.03	5.58	8.39	8.57	1.83	4.32	4.88	3.69	4.06
WT2RP3001938	0.69	0.54	7.42	4.87	0	1.52	0	1.16	0.26
WT2RP3001943	188.81	183.8	152.87	80.6	88.78	128.25	85.31	121.73	115.26
WT2RP3001944	17.12	20.99	26.58	14.69	15.73	4.83	10.12	15.18	8.75
WT2RP3001945	39.73	40.32	37.82	24.74	28.95	28.23	17.81	18.41	14.28
WT2RP3001947	47.48	38.38	48.19	30.55	15.52	22.26	24.69	18.23	25.96
WT2RP3001949	37.28	38.65	47.2	28.25	28.78	28.05	30.78	28.58	28.74
WT2RP3001952	10.17	10.07	8.89	8.88	1.57	3.07	2.44	1.18	1.17
WT2RP3001954	9.56	16.64	17.64	8.37	4.32	7.01	11.88	5.36	4.75
WT2RP3001956	3.88	4.88	8.37	7.17	1.4	2.11	0.35	2.88	1.91
WT2RP3001987	7.48	3.68	9.02	8.89	3.53	3.88	3.05	4.38	2.97
WT2RP3001989	2.15	1.05	4.78	4.72	1.42	2.2	1.83	2.58	0.82
WT2RP3001976	1.69	2.22	5.25	5.93	1.38	2.82	1.33	1.09	1.87
WT2RP3001985	3.89	3.3	6.39	4.92	1.48	2.65	3.43	6.44	2.42
WT2RP3001989	6.86	8.64	6.38	6.85	2.81	1.73	3.37	1.97	1.12
WT2RP3002002	7.76	11.38	8.67	5.38	2.01	2.22	6.25	3.04	6.18
WT2RP3002004	7.76	12.41	12.05	6.85	3.18	4.05	3.73	2.48	3.7
WT2RP3002007	14.2	10.34	19.85	11.2	4.92	6.67	4.6	3.44	6.76
WT2RP3002014	2.11	1.84	5.64	7.64	3.81	2.33	1.74	3.76	3.1
WT2RP3002015	18.24	10.71	16.79	11.55	11.36	10.60	8.87	8.67	11.36
WT2RP3002038	8	6.97	11.82	6.98	3.29	5.92	6.91	8.34	9.48
WT2RP3002045	7.45	16.7	16.88	5.97	3.53	4.5	4.78	3.11	8.05
WT2RP3002054	282.38	485.82	456.81	208.09	224.67	171.68	441.79	291.3	543.12
WT2RP3002058	3.09	3.6	6.88	3.45	0.32	0.79	0.82	1.02	0.87
WT2RP3002057	16.28	15.21	13.61	7.78	1.9	3.69	9.33	6.11	3.21
WT2RP3002061	3.7	2.82	3.37	5.14	1.22	2.6	0	1.37	0.63
WT2RP3002062	14.95	11.22	12.85	12.28	11.33	13.44	7.88	11.32	11.12
WT2RP3002063	3.07	0.91	3.18	5.78	3.45	2.63	1.35	2.97	1.84
WT2RP3002064	2.75	4.07	6.08	8.44	3.11	3.28	2.89	6.81	3.68
WT2RP3002071	4.99	2.02	2.84	4.58	1.41	2.29	2.09	2.02	0.59
WT2RP3002073	4.18	1.73	3.01	5.89	0.35	0	3.4	1.98	3.29
WT2RP3002074	3.07	4.7	9.59	5.84	1.52	1.74	2.26	2.79	1.06
WT2RP3002075	115.81	138.72	152.03	81.15	60.41	68.64	138.08	127.44	140.01
WT2RP3002077	1.88	1.37	7.12	4.73	0	2.90	1.52	0.95	1.4
WT2RP3002081	5.52	4.84	17.43	9.04	2.14	3.38	6.77	6.8	8.69

91 / 157

2621

2622

WT2RP3002089	9.1	5.64	14.68	7.72	4.13	4.75	9.35	9.69	7.69
WT2RP3002094	3.34	2.83	6.44	4.61	2.06	2.3	2.41	5.92	1.29
WT2RP3002096	12.33	8.92	11.73	8.69	3.97	5.5	5.94	5.18	4.01
WT2RP3002097	23.91	20.85	34.1	19.75	11.2	19.57	12.95	19.73	14.4
WT2RP3002098	5.03	5.84	8.49	7.72	2.61	2.62	5.48	4.78	3.63
WT2RP3002102	13.2	5.06	16.39	11.41	5.66	7.42	3.98	4.11	3.36
WT2RP3002106	10.35	7.3	14.64	10.63	8.57	6.26	3.95	4.87	10.6
WT2RP3002108	4.96	2.64	12.49	5.65	1.44	2.35	2.54	2.99	5.43
WT2RP3002109	40.69	15.02	38.55	15.39	16.89	11.31	8.11	16.62	8.14
WT2RP3002110	5.34	2.87	6.32	5.23	2.17	1.37	2.33	2.43	0.38
WT2RP3002113	9.48	7.75	15.49	10.65	5.09	3.01	6.14	4.67	6.08
WT2RP3002120	7.73	8.74	11.8	6.57	3.91	2.89	5.04	3.74	4.89
WT2RP3002121	1.11	4.94	1.77	5.17	1.32	0.68	2.8	1.37	1.2
WT2RP3002126	64.09	54.26	55.37	40.43	28.8	44.09	33.92	65.07	39.23
WT2RP3002128	8.94	4.6	9.8	7.18	4.51	3.23	4.1	5.32	11.38
WT2RP3002130	45.16	21.81	38.29	25.97	19.96	17.62	5.74	23.29	23.75
WT2RP3002133	713.78	729.07	668.65	842.53	786.78	697.72	544.62	1065.2	813.76
WT2RP3002136	99.29	50.26	79.68	69.55	58.35	39.45	17.4	37.71	62.59
WT2RP3002140	3.69	7.52	7.94	5.4	2.45	3.88	2.99	3.81	3.42
WT2RP3002142	9.07	12.94	11.88	8.33	6.6	5.19	5.4	6.06	2.64
WT2RP3002146	9.3	10.61	11.18	9.37	7.32	7.23	12.19	11.03	8.72
WT2RP3002147	11.07	11.26	19.71	10.63	7.44	5.02	4.38	6.88	4.67
WT2RP3002151	15	10.97	26.17	14.01	7.18	7.82	7.02	15.09	18.04
WT2RP3002155	53.25	64.04	47.65	73.17	43.36	52.61	36.35	94.67	86.47
WT2RP3002156	3.79	2.7	6.74	4.69	0.75	1.4	1.04	8.59	1.13
WT2RP3002160	9.08	10.26	12.47	12.87	7.96	7.04	2.86	3.65	6.87
WT2RP3002163	13.64	21.03	16.3	10.66	7.08	4.19	9.73	11.32	12.78
WT2RP3002165	6.76	7.58	8.41	6.08	1.96	1.96	2.6	3.84	1.9
WT2RP3002166	5.43	5.03	4.69	5.27	1.55	0.67	2.88	6.31	2.47
WT2RP3002173	517.9	448.69	609.69	287.97	240.8	263.29	44.88	135.22	126.7
WT2RP3002174	139.22	107.95	156.75	150.48	138.99	121.68	76.48	86.98	88.77
WT2RP3002181	5.97	4.69	12.61	5.8	2.39	3.23	1.84	5.23	1.34
WT2RP3002185	44.31	38.54	51.77	25.79	16.42	22.4	38.06	52.39	48.7
WT2RP3002193	7.33	7.63	13.67	8.26	3.71	3.99	4.19	5.78	8.23
WT2RP3002204	41.05	45.56	52.03	45.02	28.38	25.63	28.81	47.28	47.76
WT2RP3002244	12.8	20.63	18.31	13.33	15.24	10.05	9.17	7.08	5.13
WT2RP3002248	7.90	7.8	12.04	10.57	4.97	3.5	9.37	6.63	3.98
WT2RP3002252	17.21	11.16	18.79	9.56	7.33	7.05	3.72	10.64	9.39
WT2RP3002255	3.26	2.63	8.67	4.41	1.65	2.02	1.36	3.59	1.93
WT2RP3002264	81.68	74.01	84.3	98.72	70.97	43.12	54.37	98.78	53.26
WT2RP3002267	2.25	3.31	8.97	3.82	0.82	1.41	0.79	1.35	0.88
WT2RP3002273	20.4	18.49	19.5	15.02	8.97	6.71	10.11	8.27	13.28
WT2RP3002276	3.97	2.93	8.36	4.35	0.7	1.55	1.24	0.69	0.84
WT2RP3002281	13.35	12.31	15.77	11.45	7.04	7.28	10.16	10.78	11.32
WT2RP3002286	16.62	19.5	20.21	12.28	6.1	7.69	4.26	3.16	1.57
WT2RP3002297	94.9	90.81	132.4	64.72	33.84	39.98	11.41	37.45	29.98
WT2RP3002301	11.34	12.34	18.57	7.11	3.98	4.21	2.56	4.64	4.68
WT2RP3002303	7.54	10.15	15.1	7.32	5.62	4.17	5.05	6.82	7.65
WT2RP3002304	91.67	49.25	50.79	81	77.65	34.95	31.29	39.07	25.04
WT2RP3002309	4.32	3.35	10.38	4.01	1.44	1.07	2.59	1.38	0.22
WT2RP3002311	9.65	9.45	12.78	9.08	8.57	4.41	3.9	4.09	5.31
WT2RP3002316	16.31	24.1	18.79	7.98	7.42	8.29	9.67	8.19	10.65
WT2RP3002318	8.49	14.45	9.47	9.99	3.2	4.43	5.43	3.04	5.3
WT2RP3002324	18.48	20.79	21.07	16.08	8.05	11.92	2.65	8.18	8.78
WT2RP3002330	14.44	26.39	29.38	13.92	12.9	11.42	5.88	11.72	10.72
WT2RP3002333	41.51	67.04	49.57	36.81	41.81	35.64	33.68	44.76	37.87
WT2RP3002337	4.44	8.96	11.93	5.9	1.82	3.91	2.32	2.11	2.65
WT2RP3002342	25.61	22.15	24.13	19.81	11.87	9.75	11.85	13.94	20.4
WT2RP3002343	3.99	7.39	7.82	6.18	2.92	2.42	2.4	1.77	3.64
WT2RP3002351	20.98	17.51	19.58	6.79	4.97	3.6	9.93	4.83	3.54
WT2RP3002352	29.92	68.54	51.75	25.45	36.1	22.42	47.9	10.88	7.27
WT2RP3002353	28.94	28.48	26.72	23.44	18.05	15.48	14.2	29.02	22.52
WT2RP3002362	18.42	12.38	24.27	11.02	8.28	7.65	12.53	15.38	16.92
WT2RP3002383	11.71	14.02	21.35	9.9	11.24	7.29	10.05	11.93	6.97

92 / 157

2623

2624

NT2RP3002177	20.63	15.78	18.8	13.04	8.6	12.99	16.42	23.7	18.15	*	-	-
NT2RP3002187	5.32	9.77	13.16	5.81	2.98	3.19	4.39	2.85	3.83	*	-	-
NT2RP3002404	6.64	8.21	7.65	4.33	1.3	1.29	2.68	0.69	1.15	*	-	-
NT2RP3002410	7.82	4.7	8.02	3.73	3.33	1.2	3.81	3.45	2.67	*	-	-
NT2RP3002414	5.43	4.93	6.03	6.84	1.85	2.91	4.09	1.68	3.33	*	-	-
NT2RP3002430	5.86	5.44	9.44	4.78	1.52	1.85	3.2	1.23	1.16	*	-	-
NT2RP3002448	12.97	9.25	9.52	5.21	1.95	1.81	2.87	2	4.7	**	-	-
NT2RP3002454	12.06	18.92	16.37	13.74	7.64	12.02	13.75	9.17	8.67	*	-	-
NT2RP3002455	4.78	3.53	5.11	5.77	1.51	4.71	3.7	5.48	5.28	*	-	-
NT2RP3002456	58.03	38.92	43.81	27.88	14.89	15.06	14.57	38.88	31.25	*	-	-
NT2RP3002462	20.18	13.39	21.5	10.24	9.33	4.6	3.9	5.28	3.19	*	-	-
NT2RP3002468	3.83	1.84	6.7	0.93	3.37	1.82	2.45	1.7	0.82	*	-	-
NT2RP3002470	35.44	35.5	72.47	34.81	21.72	19.39	33.8	20.85	19.08	*	-	-
NT2RP3002484	3.62	2.13	4.4	4.15	1.03	1.2	2.28	1.6	1.75	*	-	-
NT2RP3002491	13.97	21.34	15.21	12.05	5.93	4.04	5.95	7.05	7.23	*	-	-
NT2RP3002494	24.08	40.78	35.02	15.6	8.77	12.21	26.54	16.61	15.03	*	-	-
NT2RP3002497	1.45	2.72	2.12	3.98	0.95	2.47	1.03	1.64	2.87	*	-	-
NT2RP3002500	18.58	14.89	20.26	24.82	15.49	27.1	14.57	11.42	19.8	*	-	-
NT2RP3002501	32.52	29.62	45.43	41.84	21.97	31.25	12.71	18.89	18.63	*	-	-
NT2RP3002512	5.61	3.7	5.13	8.12	4.35	5.63	3.81	5.5	5.08	*	-	-
NT2RP3002520	570.71	676.56	723.55	824.84	401.81	698.08	438.95	889.52	707.21	*	-	-
NT2RP3002533	33.09	38.94	36.28	44.79	15.69	31.1	9.54	21.79	17.75	**	-	-
NT2RP3002538	6.02	8.38	9.74	6.07	3.74	0.91	2.47	0	0.12	**	-	-
NT2RP3002540	4.32	7.97	5.52	9.08	2.78	5.47	9.44	4.32	9.18	*	-	-
NT2RP3002543	12.62	4.6	12.85	12.84	5.24	6.49	9.1	11.85	10.42	*	-	-
NT2RP3002546	2.89	1.16	9.33	6.12	1.48	2.25	2.07	3.55	0.82	*	-	-
NT2RP3002549	6.99	4.28	13.8	7.32	1.84	2.44	4.48	4.58	2.91	*	-	-
NT2RP3002552	4.38	5.59	10.04	7.03	1.31	2.23	3.56	3.8	2.29	*	-	-
NT2RP3002558	34.53	21.49	27.53	25.41	11.35	21.08	37	39.71	40.69	*	-	-
NT2RP3002565	13.64	7.8	14.18	11.1	3.88	9.31	11.41	13.37	18.44	*	-	-
NT2RP3002568	7.8	8.76	13.33	16.41	6.98	8.51	6.53	7.76	8.69	*	-	-
NT2RP3002571	8.13	11.03	10.93	9.57	4.89	4.39	4.63	2.87	3.9	**	-	-
NT2RP3002572	74.15	45.74	56.78	41.43	38.41	30.38	19.06	45.1	45.35	*	-	-
NT2RP3002573	32.62	15.75	29.51	19.71	16.82	14.99	13.13	12.23	21.22	*	-	-
NT2RP3002577	47.28	29.38	41.83	19.87	14.5	21	17.59	15.85	17.9	*	-	-
NT2RP3002579	8.63	8.06	12.6	7.23	3.12	4.38	5.33	4.66	5.1	*	-	-
NT2RP3002582	6.19	3.82	10.45	7.84	3.58	3.38	4.87	4.78	3.74	*	-	-
NT2RP3002587	10.07	10.45	15.78	10.84	6.77	7.46	4.8	4.74	3.88	*	-	-
NT2RP3002590	544.94	899.14	857.25	489.72	627.82	453.24	363.97	273.85	682.87	*	-	-
NT2RP3002602	7.18	7.83	10.58	10.16	7.38	6.68	4.87	4.34	8.16	*	-	-
NT2RP3002603	14.44	13.52	18.75	16.38	10.68	15.03	8.16	15.4	88.12	*	-	-
NT2RP3002621	0.81	5.15	9.83	8.58	3.04	3.97	3.12	6.17	6.09	*	-	-
NT2RP3002622	8.75	5.01	12.62	8.8	3.27	3.73	2.74	4.44	11.45	*	-	-
NT2RP3002624	26.98	16.99	21.5	17.24	10.36	13.21	17.97	25.27	18.08	*	-	-
NT2RP3002628	6.59	5.38	9.7	6.43	2.65	4.48	4.24	5.21	2.97	*	-	-
NT2RP3002629	5.76	11.73	14.59	10	6.43	5.93	6.71	4.40	3.96	*	-	-
NT2RP3002631	40.35	45.23	48.18	25.1	22.1	27.75	26.27	23.38	21.28	**	-	-
NT2RP3002647	12	15.89	13.32	16.2	15.29	15.81	10.24	7.5	7.23	*	-	-
NT2RP3002648	9.3	7.92	12.59	9.96	4.23	5.01	3.45	5.1	5.57	*	-	-
NT2RP3002650	26.86	18.53	27.47	18.25	9.88	13.45	17.35	20.46	18.82	*	-	-
NT2RP3002652	5.52	5.57	14.03	5.81	3.78	3.01	2.75	4.84	3.27	*	-	-
NT2RP3002654	20.14	13.49	14.7	11.81	10.11	6.89	5.42	7.8	5.1	*	-	-
NT2RP3002657	16.5	15.29	20.85	14.05	6.31	13.07	15.2	14.6	14.98	*	-	-
NT2RP3002659	15.92	19.34	18.86	13.66	10.7	8.74	4.85	10.09	8.73	*	-	-
NT2RP3002663	5.68	10.19	9.81	8.43	3.68	4.96	6.72	4.39	3.92	*	-	-
NT2RP3002665	10.53	17.57	11.68	14.14	15.43	9.63	19.1	11.04	8.75	*	-	-
NT2RP3002664	28.81	23.83	39.18	18.72	11.43	12.27	7.82	17.21	16.51	*	-	-
NT2RP3002667	32.83	27.15	33.62	23.84	18.84	18.81	15.18	21.9	26.19	*	-	-
NT2RP3002671	38.88	30.21	40.39	24.11	16.4	14.74	30.8	35.93	33.74	*	-	-
NT2RP3002682	8.45	6.51	13.88	7.64	3.93	4.02	2.41	3.04	2.62	*	-	-
NT2RP3002684	20.03	14.18	18.81	8.9	9.81	5.51	4.13	6.96	4.79	**	-	-
NT2RP3002687	8.25	11.21	13.71	14.12	8.33	10.78	7.88	7.74	7.01	*	-	-
NT2RP3002688	168.35	108.47	188.15	190.36	184.88	198.08	153.87	87.55	117.43	*	-	-
NT2RP3002698	8.07	8.3	9.14	8.34	5.8	6.56	5.59	6.87	4.73	**	-	-

93/157

2625

2626

NT2RP3002701	11.4	8.8	15.19	9.84	4.21	7.04	4.79	10.11	7.19				
NT2RP3002705	30.23	29.89	43.68	30.05	23.71	23.42	20.15	25.03	27.52				
NT2RP3002708	25.52	23.92	35.27	19.02	18.81	19.8	19.34	22.82	17.32				
NT2RP3002711	87.34	128.24	131.86	149.35	126.88	84.65	15.58	87.78	38.54	00	-	-	-
NT2RP3002712	11.51	9.81	14.14	7.18	5.15	2.41	5.29	7.08	4.78	0	0	-	-
NT2RP3002718	17.13	13.75	17.42	8.41	8.55	4.38	13.41	9.28	8.38	0	0	-	-
NT2RP3002721	70.08	117.72	122.3	101.88	70.22	94.12	113.7	122.48	107.48				
NT2RP3002722	20.45	20.54	15.58	16.97	11.11	10.65	14.55	10.8	10.74				
NT2RP3002723	12.21	14.76	21.14	7.75	4.55	7.78	4.05	7.72	8.91	0	0	-	-
NT2RP3002737	4.67	4.9	12.9	7.24	2.22	4.68	2.82	1.49	3.15	0	0	-	-
NT2RP3002738	122.73	88.31	164.33	158.37	134.11	152.93	81.07	85.63	82.42				
NT2RP3002742	4.82	5.31	8.08	8.59	2.59	2.11	2.29	3.26	2.78	0	-	-	-
NT2RP3002744	11.76	7.9	17.24	10.03	4.45	8.38	6.88	8.18	8.57				
NT2RP3002750	8.91	8.84	6.82	7.81	5.13	2.83	2.84	1.31	1.38	00	-	-	-
NT2RP3002757	18.22	14.88	15.85	13.82	8.41	8.83	10.99	9.29	10.38	0	00	-	-
NT2RP3002758	16.5	15.89	16.85	13.98	8.17	9.33	15.83	11.78	10.47	0	-	-	-
NT2RP3002762	6.55	8.17	14.93	9.33	2.38	7.38	4.58	5.11	8				
NT2RP3002763	6.01	4.94	14.84	7.96	3.26	4.48	4.15	2.77	3.35				
NT2RP3002770	13.79	15.86	20.19	13.26	7.67	4.65	6.88	8.79	5.16	00	-	-	-
NT2RP3002771	6.48	3.94	12.18	8.16	4.01	1.77	3.51	4.07	2.45				
NT2RP3002785	7.93	7.61	12.09	7.52	4.23	4.31	4.14	3.39	4.41	0	-	-	-
NT2RP3002789	12.74	8.68	16.54	10.03	4.49	9.87	11.67	8.3	11.43				
NT2RP3002799	44.32	36.15	61.61	71.32	40.17	61.97	38.77	29.46	32.11				
NT2RP3002801	9.13	7.95	8.02	8.82	3.17	7.55	8.99	2.43	2.83	0	-	-	-
NT2RP3002802	8.6	5.15	13.08	9.48	2.83	4.63	3.71	3.1	3.55				
NT2RP3002810	21.81	13.43	30.89	15.94	9.02	13.72	15.57	16.49	23.18				
NT2RP3002818	12.28	21.33	22.28	12.03	10.84	9.8	12.08	10.85	7.02				
NT2RP3002821	3.08	2.78	7.5	5.13	0.88	2.74	3.88	1.7	1.96				
NT2RP3002823	5.94	5.9	9.87	7.27	1.84	2.08	2.79	3.12	3.09	0	-	-	-
NT2RP3002825	46.16	88.1	89.32	16.84	18.8	11.22	9.52	7.53	4.82	00	00	-	-
NT2RP3002829	36.39	48.01	39.22	22.92	15.02	11.21	24.02	7.88	9.84	00	0	-	-
NT2RP3002831	9.11	9.95	10.41	14.39	9.89	8.12	9.98	4.98	9.9				
NT2RP3002838	16.68	11.51	17.27	10.91	4.09	9.65	4.08	4.72	7.35	00	-	-	-
NT2RP3002845	38.76	32.47	30.9	12.04	7.1	5.17	5.18	12.27	9.29	00	00	-	-
NT2RP3002852	27.07	21.18	24.92	24.24	10.22	18.18	20.47	22.49	18.87				
NT2RP3002861	6.45	10.44	9.71	7.81	2.41	4.65	6.77	4.65	5.14				
NT2RP3002869	6.19	9.1	10.7	6.8	3.23	3.7	4.89	3.27	3.99	0	-	-	-
NT2RP3002874	12.83	13.11	17.2	9.07	5.37	6.1	8.09	3.93	8.08	0	0	-	-
NT2RP3002875	143.63	148.27	181.51	294.93	163.87	216.77	137.83	99.6	141.63				
NT2RP3002877	18.65	19.83	22.97	18.33	13.19	15.19	13.02	8.47	12.29	0	-	-	-
NT2RP3002887	11.91	8.65	14.58	12.86	6.78	9.8	8.18	11.65	6.83				
NT2RP3002900	24.9	40	47.49	50.72	36.64	57.77	42.16	46.88	41.82				
NT2RP3002902	50.88	60.65	63.54	17.78	28.94	15.05	19.17	28.83	23.73	00	00	-	-
NT2RP3002909	5.43	3.6	6.73	6.91	1.07	4.91	4.39	3.43	4.07				
NT2RP3002911	32.27	38.37	39.37	37.75	22.31	32.68	41.96	41.35	47.38				
NT2RP3002948	97.24	122.64	134.73	163.88	103.17	110.88	169.55	85.09	95.2				
NT2RP3002953	130.89	155.36	174.67	145.08	99.42	98.1	201.22	181.14	165.58				
NT2RP3002955	11	13.27	12.67	9.74	5.71	6.63	7.47	5.75	5.17	0	00	-	-
NT2RP3002958	11.7	7.18	10.81	11.8	8.8	11.78	8.8	13.15	14.41				
NT2RP3002969	10.47	5.87	11.5	11.48	3.67	3.82	2.62	3.28	4.09	0	-	-	-
NT2RP3002972	9.18	3.34	9.31	7.48	3.85	0.87	5.21	5.53	7.19				
NT2RP3002976	17.8	13.14	17.07	18.61	7.47	18.55	15.56	17.59	21.58				
NT2RP3002983	14.39	18.02	14.98	15.99	8.52	11.68	18.52	16.38	19.45	0	+	-	-
NT2RP3002985	4.65	5.27	7.25	5.46	1.04	3.38	4.84	3.07	3.43				
NT2RP3002988	8.52	6.19	9.46	9	4.1	8.27	7.55	8.24	9.04				
NT2RP3003000	3.67	3.71	3.5	6.2	1.26	2.48	2.37	2.4	2.73	00	-	-	-
NT2RP3003008	10.98	2.87	10.84	10.43	9.08	7.18	5.31	8.35	5.42				
NT2RP3003012	2.33	3.47	10.69	4.62	9.4	1.73	0.84	1.68	1.54				
NT2RP3003015	94.04	42.5	71.87	52.22	37.75	51.79	66.38	44.43	68.31				
NT2RP3003018	4.14	3.97	9.63	5.88	1.6	1.48	2.55	2.43	2.28				
NT2RP3003028	3.8	4.39	9.49	7.13	4.25	1.88	9.07	2.88	1.84				
NT2RP3003029	10.59	12.64	16.85	9.48	6.29	7.18	11.38	8.45	12.28				
NT2RP3003032	18.44	18.02	27.77	16.77	11.22	12.42	15.35	17.44	10.45				
NT2RP3003041	9.55	9.82	9.95	12.39	5.68	4.67	6.47	4.02	3.62	00	-	-	-

94 / 157

2627

2628

NT2RP3001044	20.51	12.05	18.28	17.96	10.92	17.93	12.47	17.92	24.95				
NT2RP3001047	318.58	233.21	307.78	283.91	168.8	215.31	289.99	339.63	305.02				
NT2RP3001050	6.01	3.67	13.67	6.04	1.27	2.97	2.45	4.04	3.78				
NT2RP3001053	4.73	4.39	9.42	5.4	1.52	3.2	1.65	2.69	4.02				
NT2RP3001059	3.4	2.18	5.45	4.75	1.77	1.03	2.8	1.96	1.97				
NT2RP3001061	3.92	4.63	10.62	7.14	5.71	3.17	3.69	3.67	3.35				
NT2RP3001068	13.64	16.72	19.09	13.94	11.08	9.11	15.51	9.65	9.79				
NT2RP3001071	501.47	582.68	629.15	407.82	335.95	447.01	379.51	122.11	170.7	*	*	-	-
NT2RP3001076	11.62	4.82	12.85	9.77	5.93	5.76	7.19	8.4	27.45				
NT2RP3001078	14.69	14.12	17.65	8.88	7.71	7.1	3.81	7.39	11.24	**	*	-	-
NT2RP3001081	14.34	14.21	20.22	13.69	9.02	9.23	12.85	17.91	10.55				
NT2RP3001083	4.38	6.86	9.84	6.19	3.29	2.43	4.28	4.05	2.84				
NT2RP3001087	14.02	16.08	14.8	11.68	5.57	6.77	8.05	9.19	6.11	*	**	-	-
NT2RP3001088	8.1	7.48	9.23	6.48	4.67	4.01	3.89	4.01	3.83	*		-	-
NT2RP3001101	11.04	12.95	14.52	11.98	6.27	8.65	11.09	9.21	8.88				
NT2RP3001109	27.77	29.32	24.18	32.48	26.06	31.9	22.67	12.21	14.33	*		-	-
NT2RP3001121	14.74	6.75	15.15	12.65	4.86	5.96	4.11	5.09	12.75				
NT2RP3001133	8.34	7.45	12.11	8.93	5.28	6.18	2.52	4.07	7.32				
NT2RP3001137	4.53	5.11	11.1	4.41	2.55	1.77	1.57	3.58	2.61				
NT2RP3001138	14.49	14.02	31.67	15.47	8.83	7.42	8.07	8.92	9.18				
NT2RP3001150	17.72	13.28	16.68	11.32	6.97	7.87	7.6	10.95	5.4	*	*	-	-
NT2RP3001145	29.81	27.35	41.27	37.92	28.88	28.23	19.57	16.3	17.5	*		-	-
NT2RP3001160	3.07	5.2	3.99	6.23	1.85	1.66	3.17	2.48	1.85				
NT2RP3001157	36.91	28.34	31.25	37.59	28.43	24.81	16.55	12.31	12.99	**		-	-
NT2RP3001185	12.88	12.18	22.32	13.28	18.08	11.04	7.56	13.67	13.84				
NT2RP3001183	66.79	61.34	61.74	61.27	47.04	47.44	81.86	78.57	80.02				
NT2RP3001187	14.34	9.98	23.45	8.85	5.2	5.64	6.63	9.8	4.5				
NT2RP3001203	13.42	14.29	19.75	14.19	7.85	8.04	8.52	12.56	10.13				
NT2RP3001204	19.08	17.04	30.4	16.33	18.47	7.75	7.19	12.58	11.16				
NT2RP3001210	6.33	8.19	12.29	9.16	9.84	6.29	8.27	7.36	5.48				
NT2RP3001212	8.37	8.5	11.66	8.16	6.18	3.57	6.2	5.26	6.14	*		-	-
NT2RP3001213	5.67	4.05	5.85	6.17	2.96	2.21	4.78	3.58	3.94				
NT2RP3001224	55.68	53.67	56.74	16.3	18.72	16.37	11.98	29.14	28.24	**	**	-	-
NT2RP3001226	4.57	3.58	10.67	5.47	2.35	3.36	2.91	2.15	2.81				
NT2RP3001230	7.29	5.28	14.91	8.17	2.93	3.43	3.72	2.45	3.61				
NT2RP3001235	78.63	66.15	108.5	95.3	92.58	80.75	55.64	70.39	55.23				
NT2RP3001242	73.44	66.48	77.64	45.7	33.8	25.86	41.52	43.2	38.85	**	**	-	-
NT2RP3001251	104.1	73.65	128.64	135.43	95.34	99.58	79.05	81.02	71.32				
NT2RP3001262	10.37	11.27	12.13	9.17	5.92	6.54	5.47	5.61	2.73	*	**	-	-
NT2RP3001258	15.2	11.75	13.65	15.47	7.62	6.38	7.6	4.08	5.36	**		-	-
NT2RP3001260	20.33	21.23	23.55	13.01	8.75	11.13	14.88	15.72	16.47	**	**	-	-
NT2RP3001264	39.02	33.18	48.85	37.17	29.58	23.32	18.19	26.52	26.18	*		-	-
NT2RP3001273	4.51	4.91	16.07	7.81	4.38	2.69	2.42	2.45	2.99				
NT2RP3001278	10.07	13.81	16.32	10.69	5.74	4.93	4.88	5.26	3.08	**		-	-
NT2RP3001280	8.01	8.36	9.74	6.35	2.41	1.58	1.8	2.02	2.23	*	**	-	-
NT2RP3001282	6.34	4.14	6.3	6.51	1.72	3.27	3.2	1.42	0.91	*		-	-
NT2RP3001290	5.36	4.84	5.82	6.84	2.67	3.3	2.13	0.74	2.31	**		-	-
NT2RP3001301	10.24	13.13	10.58	8.04	5.22	4.9	4.68	1.19	1.93	*	**	-	-
NT2RP3001302	44.8	31.52	44.49	25.69	18.4	28.65	36.54	47.16	60.3	*		-	-
NT2RP3001311	4.75	4.55	11.5	8.43	3.14	4.48	2.2	3.71	1.64				
NT2RP3001312	24.38	35.18	29.14	9.01	9.9	12.13	5.11	5.46	5.49	**	**	-	-
NT2RP3001313	11.85	10.76	12.83	7.85	5.85	3.91	3.77	4.31	3.7	*	**	-	-
NT2RP3001327	9.48	7.49	11.81	7.32	3.83	2.99	4.81	4.57	4.3	*		-	-
NT2RP3001330	28.77	45.14	40.85	24.02	22.59	18.95	15.05	12.99	8.77	*	**	-	-
NT2RP3001344	7.42	9.51	10.17	9.58	3.28	4.53	3.31	3.28	4.42	**		-	-
NT2RP3001348	7.58	7.33	10.55	7.98	3.11	4.95	5.04	6.17	6.71				
NT2RP3001349	77.08	82.34	60.61	59.72	39.01	41.31	20.2	47.84	67.68	*		-	-
NT2RP3001353	5.79	9.35	12.49	5.84	3.68	3.54	2.78	3.58	3.64	*		-	-
NT2RP3001354	58.09	83.38	122.44	53.2	58.67	10.88	100.88	81.77	52.32				
NT2RP3001368	10.01	14.26	18.12	8.85	5.28	5.67	4.9	3.64	2.35	*	*	-	-
NT2RP3001375	13.81	11.33	19.09	11.82	5.85	11.47	12.29	9.6	11.68	*		-	-
NT2RP3001377	8.13	11.7	14	6.42	2.73	1.97	7.35	4.71	8.09	*		-	-
NT2RP3001384	15.97	18.59	24.94	10.9	5.8	5.57	11.93	4.66	4.64	*	*	-	-
NT2RP3001385	10.44	16.16	15.37	14.39	15.35	10.69	10.54	6.35	9.2				

95 / 157

2629

2630

NT2RP3003396	16.4	12.74	16.94	16.32	6.79	16.93	14.73	19.34	18.49
NT2RP3003403	3.1	2.78	7.09	5.16	1.78	3.29	1.96	3.43	3.41
NT2RP3003408	7.35	11.81	12.63	9.44	12.98	4.77	6.95	7.18	6.37
NT2RP3003411	11.08	12.25	14.60	7.42	4.83	5.8	7.12	7.74	5.44
NT2RP3003420	19.08	32.9	36.55	21.71	11.78	14.57	23.43	20.67	21.3
NT2RP3003425	6.27	6.88	8.42	6.84	2.62	3.03	4.29	2.39	4.38
NT2RP3003426	7.48	8.53	8.3	7.22	3.59	4.34	6.93	6.05	8.2
NT2RP3003427	28.71	32.12	28.4	30.38	20.64	27.6	31.57	21.68	21.07
NT2RP3003433	4.33	3.61	3.8	5.72	1.58	3.56	2.47	5.63	2.76
NT2RP3003437	8.79	2.73	6.27	7.3	2.16	3.14	2.88	3.31	4.36
NT2RP3003444	60.3	43.28	42.61	71.29	41.8	73.4	51.67	39.73	51.98
NT2RP3003455	10.05	10.38	13.48	9.15	4.03	7.98	7.2	5.64	6.66
NT2RP3003462	10.37	9.93	17.25	9.92	5.42	4.43	5.52	5.66	5.82
NT2RP3003464	14.68	13.89	13.26	12.87	7.49	6.91	18.28	16.64	18.68
NT2RP3003469	17.22	14.94	13.68	14.5	4.68	9.64	19.05	18.28	19.4
NT2RP3003473	10.99	11.83	10.53	13.09	12.46	13.39	10.9	7.59	8.71
NT2RP3003474	3.37	3.95	4.83	6.73	2.05	5.68	2.42	1.68	2.61
NT2RP3003475	25.58	23.74	27.15	19.98	21.31	0.35	11.64	10.85	19.52
NT2RP3003480	6.09	2.59	4.63	5.62	1.64	9.28	2.21	16.1	3.36
NT2RP3003491	18.8	17.66	17.3	22.68	14.42	27.63	10.64	13.19	15.87
NT2RP3003493	4.17	0.87	1.27	6.14	0.62	2.97	3.62	2.46	1.88
NT2RP3003500	5.9	4.89	5.02	7.76	1.8	6.18	6.8	6.32	6.95
NT2RP3003527	7.67	4.86	6.47	6.19	2.35	4.17	6.12	6.18	4.88
NT2RP3003532	10.61	16.22	14.72	12.47	4.81	8.89	14.42	12.45	12.16
NT2RP3003535	38.7	29.85	32.07	25.68	21.88	25.49	14.18	14.8	18.19
NT2RP3003536	5.83	2.22	11.18	6.74	1.89	4.05	4.59	4.1	3.31
NT2RP3003543	83.01	76.4	71.54	99.52	61.99	69.1	35.13	38.35	47.64
NT2RP3003549	5.05	3.61	9.39	4.68	1.49	3.34	2.95	6.92	3.73
NT2RP3003552	5.72	3.03	6.03	6.81	1.88	1.74	4.06	3.11	1.34
NT2RP3003555	7.3	5.08	11.08	9.48	2.64	6.2	0.1	6.02	7.32
NT2RP3003559	16.08	18.05	20.35	13.82	7.72	7.64	10.69	7.28	10
NT2RP3003564	21.9	11.18	19.14	32.64	13.36	24.42	5.9	11.04	13.84
NT2RP3003572	4.45	3.71	12.63	6.23	1.91	3.3	2.99	5.76	5.28
NT2RP3003576	10.13	4.39	10.89	7.2	4.48	6.35	2.89	3.48	6.89
NT2RP3003587	4.99	3.94	10.87	4.69	3.45	2.48	4.81	2.31	2.5
NT2RP3003588	8.81	6.29	8.93	6.98	6.78	3.45	8.93	5.16	5.31
NT2RP3003592	16.87	13.2	13.28	12.11	9.09	12.9	9.83	8.69	9.69
NT2RP3003593	8.95	16.67	14.21	12.38	7.8	13.05	6.6	6.07	15.06
NT2RP3003614	16.19	15.38	25.69	16.13	12.3	18.84	17.6	16.33	16.44
NT2RP3003621	23.31	9.78	21.65	19.16	8.76	10.65	10.2	15.44	18.25
NT2RP3003625	30.16	13.92	27.9	27.09	17.35	16.31	10.08	25.7	26.09
NT2RP3003627	13.85	10.22	17.95	10.88	8	10.73	7.14	16.69	27.58
NT2RP3003636	43.68	19.06	24.84	24.67	26.12	25.77	9.71	16.11	10.76
NT2RP3003642	9.56	5.41	10.21	6.42	5.7	4.71	4.31	4.84	5.36
NT2RP3003645	11.93	7.13	13.81	10.67	6.31	7.85	4.02	4.25	4.25
NT2RP3003648	6.56	6.34	9.24	8.44	4.41	4.52	4.4	2.95	3.73
NT2RP3003649	26.09	26.77	29.81	22.55	18.26	23.64	34.18	24.81	32.1
NT2RP3003650	12.81	11.48	21.64	12.68	5.46	10.24	6.28	12.27	13.31
NT2RP3003656	6.64	4.85	10.33	8.68	1.7	4.51	3.42	4.72	11.15
NT2RP3003659	9.48	6.9	15.28	7.57	6.1	6.1	4.35	6.43	5.32
NT2RP3003662	8.5	6.39	10.9	5.37	1.92	5.94	1.35	3.6	1.87
NT2RP3003664	4.6	5.97	8.27	6.78	3.03	4.25	2.14	1.93	3.82
NT2RP3003665	11.75	13.45	13.25	12.32	5.11	6.74	4.75	7.06	5.47
NT2RP3003671	70.45	72.23	46.58	69.47	88.76	53.27	45.85	13.94	22.44
NT2RP3003672	12.42	9.87	13.48	13.04	14.88	11.85	11.18	14.12	13.48
NT2RP3003673	8.49	4.06	14.72	9.56	3.98	4.91	4.97	6.88	5.68
NT2RP3003679	16.33	6.46	20.68	8.71	6.38	8.11	4.9	8.58	8.08
NT2RP3003680	13.23	16.82	26.83	9.67	11.59	10.92	6.48	10.89	9.67
NT2RP3003686	3.02	3.35	12.97	5.63	1.59	1.88	2.08	1.47	1.07
NT2RP3003689	9.23	7.33	11.82	8.17	8.77	3.83	2.32	5.97	2.24
NT2RP3003697	8.83	7.28	6.91	7.08	5.61	8.06	2.11	3.71	4.84
NT2RP3003701	9.67	9.01	11.78	7.31	4.84	5.19	3.31	2.62	4.46
NT2RP3003704	10.73	9.78	12.38	10.7	7.93	10.02	15.87	9.66	12.47
NT2RP3003714	6.67	2.51	9.58	7.86	2.41	3.23	1.8	2.83	0.8

96 / 157

2631

2632

NT2RP3003710	7.01	8.27	14.18	5.47	2.75	3.52	8.58	5.29	3.83				
NT2RP3003721	11.04	10.15	23.8	10.84	5.81	9.58	11.28	11.19	10.89				
NT2RP3003722	3.89	8.42	15.57	8.33	6.34	5.87	3.09	6.47	3.17				
NT2RP3003728	9.91	10.2	12.37	9.7	5.8	5.18	8.04	11.22	5.51				
NT2RP3003729	15.24	10.41	15.84	9.27	8.89	4.48	6.77	7.5	8.93	*			
NT2RP3003731	21.76	24.78	24.08	12.03	11.85	6.88	4.57	8.08	5.08	**	**	-	-
NT2RP3003740	6.25	8.21	13.85	9.2	5.84	6.01	5.86	5.22	4.81				
NT2RP3003744	8.07	4.78	18.45	7.1	1.15	4.58	3.94	4.58	4.9				
NT2RP3003749	12.41	10.83	19.25	6.47	8	6.02	5.17	7.75	8.42				
NT2RP3003754	6.51	6.89	16.88	6.39	6.73	2.8	13.69	10.54	7.77				
NT2RP3003759	5.04	4.82	10.4	5.36	1.64	1.95	2.96	2.71	1.17				
NT2RP3003764	37.79	25.83	53.35	42.39	40.22	42.13	25.68	27.65	28.87				
NT2RP3003768	53.84	52.07	31.23	39.67	26.34	21.22	14.6	22.88	28.5	*			
NT2RP3003767	8.01	8.08	7.88	8.43	4.34	4.24	4.39	2.88	4.34	**			
NT2RP3003778	24.16	27.98	20.47	21.14	16.84	15.25	32.15	22.31	26.7				
NT2RP3003778	12.68	5.29	15.64	10.68	4.35	10.53	3.1	5.85	4.41				
NT2RP3003783	21.6	12.18	19.59	7.99	4.9	7.08	8.28	8.7	5.9	*	*	-	-
NT2RP3003787	6.7	5.75	13.41	6.04	3.64	1.85	4.2	4.99	1.48				
NT2RP3003789	8.11	11.38	13.5	7.92	6.55	6.87	5.52	5.69	3.28	*			
NT2RP3003795	28.23	17.85	14.78	13.94	17.1	12.18	12.09	11.24	13.99				
NT2RP3003798	7.73	3.87	9.64	5.67	2.6	3.17	2.83	1.93	2.55				
NT2RP3003800	80.77	78.75	88.8	62.18	49.84	73.93	101.81	58.83	58.22				
NT2RP3003805	19.72	20.08	15.07	17.29	8.51	14.67	17.48	12.39	11.94				
NT2RP3003809	8.63	14.84	28.37	7.81	4.06	5.88	6.78	7.04	6.7				
NT2RP3003819	19.91	12.71	21.8	8.29	12.89	15.12	8.35	11.58	14.32				
NT2RP3003824	4.44	5.34	16.42	6	3.28	2.18	5.16	4	3.14				
NT2RP3003825	15.83	8.99	17.57	10.52	9.35	8.99	8.88	7.14	8.67				
NT2RP3003828	7.18	6.86	13.27	7.4	5.95	4.81	5.75	3.03	3.24				
NT2RP3003831	49.77	38.05	46.92	12.19	13.59	13	11.7	8.2	8.68	**	**	-	-
NT2RP3003833	7	12.76	5.52	6.51	1.52	3.02	3.01	1.45	1.88				
NT2RP3003835	22.82	45.16	37.16	16	14.78	18.49	26.73	7.91	11.58	*			
NT2RP3003842	3.09	1.45	4.55	4.68	0.91	2.25	1.5	2.49	0.97				
NT2RP3003843	23.17	13.64	18.94	21.62	11.25	18.44	16.81	22.17	19.21				
NT2RP3003844	11.64	8.51	12	9.58	4.9	7.1	7.94	10.02	13.51				
NT2RP3003846	11.58	8.88	8.25	8.42	3.89	7.04	5.15	6.21	10.88				
NT2RP3003849	5.16	5.81	8.2	5.18	1.5	2.43	2.36	2.09	1.87	*			
NT2RP3003862	8.43	6.43	11.03	8.94	2.53	2.38	2.84	2.47	3.72	*			
NT2RP3003870	9.39	14.91	15.8	7.42	3.58	2.51	4.29	2.22	5.4	*	*	-	-
NT2RP3003874	8.27	8.08	7.54	6.09	3.19	3.65	2.59	2.64	5.21	*	**	-	-
NT2RP3003875	6.84	5.28	7.88	10.08	2.62	6.72	2.11	6.22	4.3				
NT2RP3003880	11.2	8.02	13.91	8.47	2.77	7.12	3.7	8.83	6.95				
NT2RP3003889	6.9	11.63	11.35	8.75	7.41	9.61	7.37	8.08	5.22				
NT2RP3003891	13.51	11.29	17.94	8.42	7.04	7.49	10.43	12.21	12.8	*			
NT2RP3003914	2.72	1.62	6.58	5.33	2.56	2.54	2.7	1.4	2.69				
NT2RP3003915	13.41	9.97	16.84	9.16	3.98	6.29	7.39	4.87	7.38	*	*	-	-
NT2RP3003918	7.35	6.41	5.89	8.06	2.84	2.83	4.28	2.94	4.43	*			
NT2RP3003920	10.75	6.92	10.79	7.41	1.85	6.22	5.71	4.38	3.03	*			
NT2RP3003924	33.62	17.52	18.55	12.63	9.6	13.46	9.41	16.84	13.76	*			
NT2RP3003932	13.16	5.83	7.58	6.21	2.22	4.38	1.9	2.22	2.59	*			
NT2RP3003939	11.85	14.78	16.54	9.97	7.98	8.34	3.34	3.94	3.84	*	**	-	-
NT2RP3003940	12.22	9.19	12.29	11.49	8.58	4.42	7.78	12.51	10.79				
NT2RP3003943	7.2	4.68	6.57	7.12	5.58	5.18	4.42	3.92	6.05				
NT2RP3003959	7.72	13.55	11.27	9.14	1.71	5.64	6.69	7.88	9.75				
NT2RP3003981	75.69	75.77	72.03	129.26	82.85	106.1	81.78	51.98	65.78				
NT2RP3003985	18.74	14.24	13.95	14.48	8.03	12.62	20.87	19.58	26.47				
NT2RP3003972	4.57	5.37	14.74	8.17	2.14	5.45	5	3.91	5.17				
NT2RP3003973	6.48	6.48	12.61	7.2	3.25	3.55	3.26	4.28	6.78				
NT2RP3003978	4.45	3.72	9.63	4.82	2.41	2.62	1.15	1.72	1.58				
NT2RP3003980	146.05	117.79	158.74	103.45	88.31	102.32	155.94	132.47	147.8	*			
NT2RP3003982	7.33	5.76	16.01	8.18	2.48	3	4.81	2.79	5.02				
NT2RP3003983	10.71	10.02	27.87	8.13	4.12	8.61	5.84	9.52	13.58				
NT2RP3003982	46.04	43.58	55.68	31.82	22.83	38.4	32.87	30.71	18.85	*	*	-	-
NT2RP3004000	75.38	38.37	57.12	93.25	54.67	51.8	38.78	36.39	33.4				
NT2RP3004001	14.94	10.18	22.03	15.95	8.73	15.13	13.18	15.23	21.08				

97 / 157

2633

2634

NT2RP3004005	23.75	15.81	26.46	14	10.84	14.12	18.28	18.58	18.43				
NT2RP3004013	22.54	11.33	21.98	5.11	3.13	3.73	3.05	4.84	3.43	*	*	-	-
NT2RP3004018	7.39	7.18	13.3	10.82	9.82	5.35	6.27	5.44	4.76				
NT2RP3004025	22.82	15.93	25.25	22.37	10.03	12.12	21.2	17.81	21.7				
NT2RP3004030	4.66	4.65	8.61	4.83	3.44	3.8	4.38	2.41	1.96				
NT2RP3004041	4.1	2.26	3.06	5.03	2.28	1.57	2.62	0.7	0.71				
NT2RP3004042	2602.1	1724.7	1778.2	2098.3	1690	1795.1	1947.3	2828.9	2085.3				
NT2RP3004044	32.34	21.11	30.01	12.91	11.29	11.35	7.18	15.29	26.2	**		-	-
NT2RP3004051	19.22	22.27	14.08	31.29	21.4	25.83	12.31	39.18	27.89				
NT2RP3004052	20.9	17.89	21.89	18.39	15.97	16.97	13.27	17.39	19.16				
NT2RP3004053	19.14	11.95	20.87	15.13	10.18	12.84	3.1	7.99	5.77	*		-	-
NT2RP3004055	5.06	6.81	8.8	5.35	3.84	2.7	2.8	3.03	2.13	*		-	-
NT2RP3004059	28.03	28.01	36.98	19.3	19.54	21.1	20.16	18	15.83	*	*	-	-
NT2RP3004083	29.14	22.42	34.14	23.83	23.28	19.51	31.01	26.64	12.72				
NT2RP3004087	7.95	8.82	20.78	10	4.39	4.65	3.75	4.12	8.39				
NT2RP3004070	905.81	934.59	985.28	291.19	822.85	595.78	278.31	1360.6	1278.5	*		-	-
NT2RP3004075	11.59	15.27	28.01	8.7	5.82	8.89	3.95	5.77	5.73	*		-	-
NT2RP3004078	13.83	11.53	17.13	13.53	5.08	9.69	12.68	11.48	12.45				
NT2RP3004083	33.39	27.25	49.84	12.1	13.49	7.53	8.88	6.07	6.78	*	**	-	-
NT2RP3004084	25.34	24.44	39.2	27	15.25	16.2	38.87	47.1	47.33				
NT2RP3004087	13.47	17.08	17.12	15.44	8.85	8.53	10.22	10.37	8.13	*		-	-
NT2RP3004090	72.24	81.09	78.4	57.09	57.97	55.38	77.78	70.88	90.88	**		-	-
NT2RP3004093	28.49	18.68	45.95	14.41	12.09	18.33	8.92	23.19	25.5				
NT2RP3004095	33.26	26.68	48.11	23.18	21.66	31.49	19.24	23.44	21.47				
NT2RP3004102	34.73	35.32	40.77	25.98	17.78	24.22	15.74	23.36	12.18	*	**	-	-
NT2RP3004110	7.05	4.57	10.07	7.67	1.49	3.89	2.25	3.65	1.82	*		-	-
NT2RP3004119	11.05	7.86	13.75	6.77	3.42	4.11	3.84	3.73	1.51	*	*	-	-
NT2RP3004125	3.06	3.32	9.52	6.05	1.93	1.78	1.81	1.49	1				
NT2RP3004129	11.05	13	16.54	13.66	11.08	8.18	7.67	3.97	5.87	*		-	-
NT2RP3004130	6.85	5.7	8.22	6.11	4.3	3.04	1.91	3.1	2.41	**		-	-
NT2RP3004133	8.31	5.24	20.22	8.3	3.21	4.89	3.33	4.08	3.29				
NT2RP3004145	5.99	4.29	13.18	5.33	2.78	2.89	2.63	4.11	3.14				
NT2RP3004148	53	37.5	49.68	32.7	21.92	15.17	10.82	36.4	44.13	*		-	-
NT2RP3004155	8.18	8.86	12.7	12.14	3.98	4.25	2.59	2.94	3.13	**		-	-
NT2RP3004185	77.97	53.41	68.82	61.37	48.12	31.89	48.51	25.55	55.19				
NT2RP3004179	9.55	6.18	11.2	8.95	2.94	2.88	4.01	3	3.7	*		-	-
NT2RP3004185	8.31	5.21	6.12	8.04	2.95	2.38	1.55	0.61	2.3	**		-	-
NT2RP3004188	8.31	7.01	6.85	7.94	4.08	2.67	3.74	2.3	1.82	**		-	-
NT2RP3004189	15.63	7.3	21.15	9.68	5.21	8.89	4.83	8.4	6.14				
NT2RP3004180	64.52	34.38	57.67	48.45	40.38	53.55	47.37	42.21	45.49				
NT2RP3004181	122.58	123.5	136.79	74.78	48.88	44.19	48.51	70.16	47.4	**	**	-	-
NT2RP3004202	18.55	13.31	21.41	14.01	15.47	6.42	10.97	12.13	10.67				
NT2RP3004205	10.59	7.33	14.79	6.1	4.73	2.82	2.52	2.35	2.6	*		-	-
NT2RP3004206	15.55	12.16	17.84	7.47	8.84	7.63	5.69	7.13	12.18	*		-	-
NT2RP3004207	8.85	8.42	9.24	5.53	2.88	3.67	2.31	1.22	2.02	**	**	-	-
NT2RP3004209	27.4	23.65	14.82	17	13.67	16.8	19.95	17.39	12.49				
NT2RP3004215	22.45	23.53	27.88	11	7.75	10.82	4.22	8.07	7.54	**	**	-	-
NT2RP3004219	5.58	5.12	13.51	6.61	2.08	4.81	3.19	3.17	3.09				
NT2RP3004242	4.18	6.25	13.02	6.15	3.05	4.41	4.72	3.98	3.45				
NT2RP3004246	3.85	4.1	9.37	4.89	2.24	1.94	1.48	3.04	2.11				
NT2RP3004253	7.46	3.55	9.97	5.38	2.59	1.44	2.57	2.47	3.21				
NT2RP3004256	8.43	4.33	9.8	7.18	4.84	3.38	5.14	4.78	3.41				
NT2RP3004262	5.75	6.49	8.12	6.38	2.36	2.74	4.69	0.6	2.65				
NT2RP3004275	5.37	6.56	6.47	6.33	1.33	2.3	2.5	1.89	3.75	**		-	-
NT2RP3004282	4.88	27.86	18.83	6.38	3.74	6.05	2.71	3.54	3.27				
NT2RP3004289	6.43	5.13	10.82	5.92	2.79	3.35	2.25	2.59	3.21				
NT2RP3004294	4.71	4.44	6.56	4.85	2.62	3.32	2.88	6.45	3.88				
NT2RP3004298	64.72	65.93	73.74	63.66	58.3	70.43	70.1	51.48	68.44				
NT2RP3004309	10	7.69	11.87	7.21	2.5	3.75	5.84	3.48	5.38	*	*	-	-
NT2RP3004321	23.29	38.14	42.57	12.44	15.53	8.78	14.14	5.43	9.71	*	*	-	-
NT2RP3004322	4.18	4.29	4.98	4.8	1.3	0.91	2.93	1.84	2.18	**		-	-
NT2RP3004332	9.53	11.83	11.13	7.92	3.85	4.05	4.3	2.24	2.92	*	**	-	-
NT2RP3004334	14.9	13.13	17.93	15.29	11.08	7.25	6.62	11.71	9.43	*		-	-
NT2RP3004336	9.94	8.5	13.65	9.26	7.37	6.68	4.73	8.17	4.32	*		-	-

98 / 157

2635

2636

NT2RP3004334	20.15	18.11	15.61	11.35	11.45	18.93	14.61	18.8	18.46		
NT2RP3004341	45.52	46.06	58.9	30.5	32.88	24.48	35.78	44.1	42.23	*	-
NT2RP3004345	5.41	7.27	12.03	7.28	2.28	2.19	5.01	3.58	4.59		
NT2RP3004348	8.2	11.9	19.15	6.37	5.32	6.29	9.92	4.97	8.08		
NT2RP3004349	46.27	44.89	52.77	25.21	14.4	28.82	51.32	39.58	50.42	**	-
NT2RP3004355	13.69	17.95	13.45	12.62	8.18	11.72	10.41	5.58	7.23		
NT2RP3004356	6.56	4.92	9.78	7.48	3.18	5.93	2.87	8.09	3.61		
NT2RP3004360	11.41	7.43	12.2	11.26	4.38	7.34	4.52	4.39	7.23	*	-
NT2RP3004361	22.43	16.96	27.29	12.91	14.7	10.73	18.55	21.02	28.27		
NT2RP3004374	6.08	4.11	5.92	5.52	2.91	4.87	2.31	2.56	2.6	*	-
NT2RP3004378	60.03	59.45	48.92	59.78	39.18	55.24	61.97	46.48	63.83		
NT2RP3004399	25.24	30.84	27.39	23.99	13.72	17.1	23.05	18.32	24.46	*	-
NT2RP3004405	6.48	2.89	3.95	8.86	3.8	2.54	2.58	3.57	3.49		
NT2RP3004408	48.02	50.1	76.08	27.28	18.49	29.22	40.17	40.49	25.13	*	*
NT2RP3004411	21.84	13.45	15.39	11.2	8.65	12.78	6.83	10.8	17.8		
NT2RP3004424	12.83	9.11	14	10.53	6.29	12.31	8.49	6.88	11.07		
NT2RP3004428	9.06	4.59	7.61	7.92	3	6.19	2.13	2.45	2.73	*	-
NT2RP3004432	41.77	40.01	49.18	48.87	29.7	37.75	43.11	55.91	45.48		
NT2RP3004434	8.59	6.38	9.2	18.42	18.09	11.41	8.35	8.97	10.87		
NT2RP3004446	5.55	1.31	2.23	6.8	1.27	2.52	2.84	1.81	2.66		
NT2RP3004451	28.05	27.38	35.9	43.8	34.57	29.99	27.84	20.66	27.08		
NT2RP3004454	3.04	4.85	3.07	5.17	2.65	2.87	3.32	1.95	4.78		
NT2RP3004468	17.88	12.48	17.83	17.19	13.78	19.32	8.38	10.38	11.73	*	-
NT2RP3004470	6.58	4.48	12.1	8.37	2.81	3.83	6.65	6.78	5.57		
NT2RP3004472	4.96	8.32	13.71	7.25	6.54	4.58	2.83	3.65	3.08		
NT2RP3004475	9.27	6	11.18	8.68	2.73	3.48	3.79	5.2	3.34	*	-
NT2RP3004480	22.33	15.69	32.46	17.18	10.83	15.48	12.05	10.67	10.96		
NT2RP3004481	7.61	6.38	10.91	7.98	3.04	8.85	8.17	5.21	5.83		
NT2RP3004490	1.96	3.67	4.31	5.07	2.67	3.48	2.86	2.05	1.78		
NT2RP3004495	28.05	43.68	52.3	32.37	24.7	28.45	17.59	12.79	20.95	*	-
NT2RP3004498	6.1	6.29	10.2	8.29	5.15	5.43	3.38	4.35	5.77		
NT2RP3004503	13.34	8.17	15.88	13.28	6.98	13.95	9.69	13.04	18.21		
NT2RP3004504	26.88	13.87	24.17	18.13	11.34	13.03	8.65	9.71	9.3	*	-
NT2RP3004505	60.58	42.22	65.16	60.93	61.18	59.83	38.93	40.66	35.92	**	-
NT2RP3004507	6.96	5.95	8.15	6.81	3.02	3.27	2.74	2.78	2.24		
NT2RP3004519	3.09	3.32	8.34	4.29	1.92	0.77	1.78	0.64	1.58		
NT2RP3004524	22.27	16.91	24.48	12.8	14.38	19.78	10.59	7.77	17.02	*	-
NT2RP3004527	31.26	36.6	41.52	32.19	22.11	22.1	23.08	21.05	18.57	**	-
NT2RP3004534	1954.9	1225.4	1438.4	1628.7	1505.7	1779.6	777.34	2507.5	2418.6		
NT2RP3004539	4.71	4.69	10.04	5.61	3.18	4.71	4.15	5.62	6.25		
NT2RP3004541	14.72	10.14	22.05	12.49	7.25	9.05	8.78	11	10.96		
NT2RP3004544	17.32	13.37	16.21	16.43	8.76	13.78	6.55	10.61	10.68	*	-
NT2RP3004551	12.43	7.15	13.73	9.94	5.53	8.65	10.18	9.42	7.62		
NT2RP3004552	18.37	11.88	21.3	18.2	12.57	18.88	18.98	16.22	18.63		
NT2RP3004557	11.18	12.99	15.15	12.15	8.14	12.54	12.08	12.56	14.07		
NT2RP3004581	2.62	4.37	4.68	6.9	2.44	3.04	3.97	3.31	3.2		
NT2RP3004586	16.46	13.14	29	10.83	7.99	8.15	3.35	6.43	21.27		
NT2RP3004589	20.41	16.28	28.7	14.13	11.71	13.43	13.97	16.78	16.76		
NT2RP3004572	12.48	13.38	19.74	11.19	9.39	9.67	3.98	9.35	8.35	*	-
NT2RP3004578	8.49	6.06	8.08	7.21	4.59	4.3	5.13	4.76	4.1	*	-
NT2RP3004584	8.34	8.1	9.67	5.07	2.29	4.8	3.85	3.5	2.08	*	-
NT2RP3004588	4.64	4.62	7.48	7.78	2.88	3.22	3.01	5.54	3.28		
NT2RP3004594	18.77	25.61	21.44	21.59	19.15	22.94	29.88	27.19	18.84		
NT2RP3004803	4.78	6.26	5.77	8.8	3.9	4.9	5.48	7.72	5.82		
NT2RP3004812	6.48	6.92	12.56	12.71	4.69	5.32	2.31	10.42	14.75		
NT2RP3004817	8.05	8.14	15.39	8.87	3.8	5.79	3.78	8.88	7.16		
NT2RP3004818	38.58	23.11	37.29	24.87	25.7	22.47	22.39	33.72	29.81		
NT2RP3004825	19.69	16.42	18.21	11.78	10.81	14.71	6.25	8.78	11.12	*	-
NT2RP3004835	11.5	9.72	13.73	7.06	5.4	4.21	3.58	2.39	2.64	*	-
NT2RP3004840	4.25	10.31	12.87	7.49	7.74	5.34	4.28	3.2	9.33		
NT2RP3004842	15.02	23.4	27.93	15.98	15.82	18.39	17.49	17.88	14.1		
NT2RP3004847	10.62	11.7	14.69	13.69	6.27	6.34	18.22	14.88	18.18	*	+
NT2RP3004852	28.1	24.38	41.78	13.95	14.47	12.02	4.02	12.27	12.85	*	-
NT2RP3004869	7.74	7.41	20.24	7.31	6.89	6.49	4.02	6.68	4.21		

99 / 157

2637

2638

NT2RP3004470	5.78	5.9	16.65	6.95	8.77	4.78	1.73	4.19	3.14				
NT2RP4000008	4.28	3.75	9.74	7.01	3.13	2.8	0.87	1.61	1.16				
NT2RP4000018	777.26	720.79	779.71	458.85	552.02	180.81	868.14	841.39	673.73	*	*	-	-
NT2RP4000023	12.08	13.49	11.95	6.59	4.27	3.93	5.43	2.46	2.58	**	**	-	-
NT2RP4000025	10.92	10.01	10.19	8.87	3.46	6.52	5.55	3.38	7.2	*	*	-	-
NT2RP4000035	18.64	12.06	18.1	13.28	10.2	10.62	12.01	14.28	15.16				
NT2RP4000041	7.95	4.85	12.17	6.47	2.4	5.6	2.83	2.65	3.55				
NT2RP4000049	5.5	8.69	14.31	6.89	4.49	2.04	3	3.25	4.73				
NT2RP4000050	7.92	6.2	13.4	9.91	4.4	3.61	2.52	5.2	2.39				
NT2RP4000051	11.64	12.9	17.08	9.37	6.75	7.7	7.83	11.79	8.18	*	*	-	-
NT2RP4000063	8.36	7.43	9.4	6.21	2.66	4.7	2.12	2.88	2.29	*	**	-	-
NT2RP4000065	8.57	5.45	7.62	5.43	1.38	3.74	3.79	2.44	1.14	*	*	-	-
NT2RP4000070	8.9	10.6	11.32	10.92	5.17	4.74	3.9	3	5.41	**	**	-	-
NT2RP4000074	7.09	6.56	7.33	6.07	3.64	2.14	2.7	1.95	3.11	**	*	-	-
NT2RP4000078	10.64	7.98	15.22	8.04	4.02	9.72	4.12	7.85	6.44				
NT2RP4000080	39.53	29.48	35.07	14.73	7.78	7.51	7.13	12.84	8.56	**	**	-	-
NT2RP4000099	17.27	15.29	21.59	9.54	12.81	8.62	5.87	6.72	3.64	*	**	-	-
NT2RP4000102	8.52	8.93	16.05	7.45	6.98	6.31	6.6	5.59	5.12				
NT2RP4000103	20.66	29.01	31.17	15.7	18.08	13.47	20.04	13.68	12.68	*	*	-	-
NT2RP4000108	10.35	7.66	11.54	5.11	6.39	6.22	4.98	2.69	3.44	*	**	-	-
NT2RP4000109	9.73	11.84	8.78	8.41	4.9	6.21	8.7	3.67	4.98	*	*	-	-
NT2RP4000111	9.68	7.22	7.83	10.68	5.18	7.24	8.51	4.5	2.88				
NT2RP4000112	16.81	11.4	24.28	20.79	18.36	22.3	16.09	13.49	17.44				
NT2RP4000115	7.34	4.25	10.88	5.53	2.81	4.32	2.76	2.51	2.27				
NT2RP4000129	24.84	25.48	33.51	13.19	12.49	13.21	12.38	15.5	12.9	**	**	-	-
NT2RP4000137	20.7	18.54	14.05	10.11	6.13	6.89	9.81	6.38	10.47	*	*	-	-
NT2RP4000138	7.91	6.25	8.72	5.36	2.95	3.73	3.09	3.11	5.54	*	*	-	-
NT2RP4000141	11.63	7.1	7.39	5.05	2.2	2.55	3.83	2.34	1.67	*	*	-	-
NT2RP4000147	7.8	8.75	6.46	5.57	2.45	1.71	3.71	1.66	2.95	*	**	-	-
NT2RP4000150	22.83	40.43	32.64	28.18	18.45	19.68	24.1	13.63	20.82				
NT2RP4000151	5.35	2.77	7.82	5.7	2.57	4.58	2.81	1.73	3.64				
NT2RP4000157	5.03	3.17	7.55	5.12	2.41	3.03	2.61	4.59	2.88				
NT2RP4000159	7.7	6.21	14	7.81	5.43	9.16	5.02	7.93	7.15				
NT2RP4000163	11.36	16.35	22.24	7.82	6.7	6.1	11.58	8.66	14.17	*	*	-	-
NT2RP4000167	5.72	10.78	11.28	8.86	1.93	2.94	6.09	1.64	2.88				
NT2RP4000171	17.67	20.8	18.76	13.24	7.81	11.48	17.22	11.6	13.85	*	*	-	-
NT2RP4000175	7.68	11.5	9.21	7.19	5.94	7.44	5.84	4.88	6.71	*	*	-	-
NT2RP4000180	8.25	9.87	9.45	8.6	4.86	7.21	9.47	6.55	7.72				
NT2RP4000185	3.44	9.69	9.07	5.25	3.82	5.08	3.89	5.51	8.51				
NT2RP4000192	3.68	2.3	7.85	7.16	4.58	3.22	2.99	2.89	2.84				
NT2RP4000194	4.94	3.92	6.27	5.67	6.92	4.33	4.73	3.32	6.3				
NT2RP4000196	4.44	3.86	8.22	4.42	2.19	1.89	2.14	2.44	1.01				
NT2RP4000210	4.32	3.8	5.46	5.03	2.17	1.97	3.56	2.52	3.91				
NT2RP4000212	10.41	10.12	9.74	6.43	6.54	3.07	4.88	4.69	2.44	*	**	-	-
NT2RP4000214	21.41	28.56	31.47	14.93	12.48	12.75	14.71	10.63	11	*	**	-	-
NT2RP4000218	18.58	17.54	17.8	9.05	3.32	6.3	7.12	4.66	4.69	**	**	-	-
NT2RP4000218	20.76	13.22	21.73	16.85	12.17	21.91	12.18	18.58	23.33				
NT2RP4000223	11.85	6.6	8.31	9.5	4.18	9.75	8.04	7.05	11.94				
NT2RP4000243	10.56	9.93	12.85	11.2	9.88	11.7	8.64	8.69	10.23				
NT2RP4000248	7.12	5.86	6.44	5.65	4.43	4.42	2.97	2.31	3.27	**	*	-	-
NT2RP4000250	11.29	8.98	12.39	10.04	3.49	5.45	5.45	5.81	5.67	**	*	-	-
NT2RP4000256	3.28	5.13	8.67	4.51	2.18	2.58	3.11	2.41	2.48				
NT2RP4000257	23.54	23.13	30.8	34.24	27.3	37.51	26.44	19.58	27.22				
NT2RP4000259	5.19	5.55	3.54	8.67	2.13	4.17	4.34	3.84	5.01				
NT2RP4000261	4.72	2.05	7.37	6.8	1.48	3.78	2.37	3.23	3.58				
NT2RP4000262	2.93	2.46	11.12	5.21	1.63	2.61	2.49	2.62	2.71				
NT2RP4000263	8.03	5.82	13.8	7.33	4.75	4.78	8.85	4.63	5.93				
NT2RP4000280	36.69	25.69	34.43	21.81	13.93	11.69	23.3	25.5	21.75	*	*	-	-
NT2RP4000285	7.96	5.71	10.34	8.48	4.4	5.47	4.58	4.72	2.93				
NT2RP4000290	8.28	5.27	10.35	7.67	3.77	4.58	5.45	2.24	4.35				
NT2RP4000291	23.37	31.04	45.88	29.65	20.44	28.9	16.58	13.08	6.43	*	*	-	-
NT2RP4000301	9.08	21.31	16.13	18.84	21.42	12.52	13.04	8.64	8.68				
NT2RP4000312	7.37	5.88	10.71	8.75	4.45	5.45	4.08	4.23	5.32				
NT2RP4000321	14.11	6.64	13.51	10.77	5.65	7.11	6.38	8.75	12.78				

100/157

2639

2640

NT2RP4000323	42.99	31.25	39.68	23.73	23.52	24.08	25.22	26.31	20.02	*	*	-
NT2RP4000324	28.87	20.72	25.53	23.74	16.73	21.66	28.32	24.57	21.68			-
NT2RP4000334	11.51	10.77	15.15	9.55	9.47	9.13	8.39	8.12	5.89	**		-
NT2RP4000343	29.88	12.62	32.18	18.49	14.22	8.72	14.98	17.89	14.7			-
NT2RP4000348	12.35	14.21	17.95	11.42	8.77	9.63	12.1	7.85	8.02			-
NT2RP4000349	8.77	18.24	12.89	11.44	12.7	9.21	11.37	7.34	11.23			-
NT2RP4000355	253.25	117.6	189.34	264.37	149.78	254.47	257.02	316.08	192.79			-
NT2RP4000356	61.98	38.7	79.81	60.03	57.95	82.79	39.95	40.28	37.12			-
NT2RP4000360	38.52	30.1	36.81	28.77	38.27	34.85	22.5	28.04	23	*		-
NT2RP4000367	5.32	7.72	13.11	9.15	7.75	8.68	6.85	7.52	4.05			-
NT2RP4000370	6.13	3.8	7.29	7.4	4.81	4.6	4.29	3.27	2.44			-
NT2RP4000373	52.18	35.58	49.83	33.3	22.86	20.18	8.95	13.1	7.55	*	**	-
NT2RP4000376	4.35	4.78	3.85	5.98	2.65	3.6	3.64	2.24	1.76	*		-
NT2RP4000381	4.21	7.62	7.46	8.53	4.41	5.48	5.18	4.44	4.11			-
NT2RP4000388	15.3	8.64	28.13	9.56	8.37	8.78	3.98	7.28	10.59			-
NT2RP4000390	9.32	12.81	21.1	9.57	7.3	8.9	4.45	8	7.73			-
NT2RP4000393	4.73	6.04	14.2	5.91	3.78	3.4	2.49	4.28	1.81			-
NT2RP4000398	15.1	12.32	17.4	15.43	7.99	12.09	10.09	7.98	7.95	*		-
NT2RP4000406	6.93	7.1	10	7.22	4.81	7.48	4.73	7.43	4.16			-
NT2RP4000407	14.2	8.55	18.88	9.39	7.38	7.23	8.05	7.66	8.08			-
NT2RP4000413	6.36	8.44	8.74	7.32	6.23	4.25	14.9	11.84	8.91			-
NT2RP4000415	5.58	6.36	6.25	8.19	3.66	6.69	6.61	5.43	4.11			-
NT2RP4000417	4.81	3.78	14.82	6.27	2.89	2.95	1.78	3.56	2.97			-
NT2RP4000423	8.58	11.85	52.11	7.59	6.28	6.14	5.59	8.25	4.46			-
NT2RP4000424	5.63	6.45	12.56	7.04	3.08	2.9	3.68	4.14	1.14			-
NT2RP4000447	7.22	11.02	14.44	6.8	2.54	3.42	2.02	2.94	1.88	*		-
NT2RP4000448	10.44	4.78	12.49	7.44	5.19	4.9	2.34	3.11	4.18			-
NT2RP4000449	14.23	13.95	15.79	8.35	8.81	8.15	8.09	11.82	7.03	**	*	-
NT2RP4000453	9.01	8.07	14.02	7.74	6.15	4.89	6.58	4.27	8.08			-
NT2RP4000455	9.03	10.34	8.68	7.99	5.45	3.17	4.44	4.36	4.13	**		-
NT2RP4000456	7.04	6.17	17.02	5.76	3.52	4.72	1.68	2.58	3.45			-
NT2RP4000457	13.26	8.71	17.6	8.13	4.23	7.52	11.15	9.14	8.58			-
NT2RP4000461	55.44	37.43	48.48	34.13	24.48	24.96	29.63	26.44	29.5	*	*	-
NT2RP4000462	42.85	30.75	56.94	75.69	50.54	40.64	52.88	43.57	42.53			-
NT2RP4000483	85.58	62.21	92.77	65.01	62.99	43.88	67.48	75.6	83.4			-
NT2RP4000471	25.28	34.12	24.81	15.24	25.16	15.87	19.8	18.58	11.2	*		-
NT2RP4000472	7.71	9.85	9.52	10.37	5.54	5.62	6.84	6.87	3.82			-
NT2RP4000476	14.81	18.77	18.55	11.79	4.98	3.82	5.79	4.35	2.67	*	**	-
NT2RP4000480	12.42	19.39	22.37	9.71	8.11	7.69	2.98	6.14	7.56			-
NT2RP4000481	7.17	5.4	16.93	8.71	1.97	4.18	2.54	3.59	3.12			-
NT2RP4000483	21.87	12.5	29.34	14.63	7.41	9.82	17.08	21.67	14.93			-
NT2RP4000487	11.87	8.84	15.81	9.26	6.43	6.73	8.08	10.31	7.08			-
NT2RP4000496	18.33	41.67	19.05	11.91	7.25	5.15	7.41	10.6	17.63			-
NT2RP4000497	5.81	3.81	10.08	3.87	3.09	2.81	2.48	1.77	2.22			-
NT2RP4000498	25.64	11.6	45.3	21.94	11.58	20.83	11.71	6.48	8.74			-
NT2RP4000500	16.18	14.2	13.24	21.4	11.25	11.2	15.02	6.9	5.02			-
NT2RP4000507	14.47	9.74	21.49	11.59	12.91	17.98	12.44	18.01	18.95			-
NT2RP4000515	34.88	28.88	36.21	23.73	25.4	27.8	24.2	28.54	29.19	*		-
NT2RP4000518	8.8	4.78	15.95	7.22	3.73	6.28	5.58	4.23	5.23			-
NT2RP4000517	6.83	4.39	11.92	5.6	2.28	2.95	2.28	2.51	3.47			-
NT2RP4000518	5.88	5.23	9.62	6.13	3.29	2.14	2.68	2.57	2.25	*		-
NT2RP4000519	11.25	13.6	9.43	7.22	5.12	6.11	9.98	4.77	2.74	*		-
NT2RP4000524	3.66	2.61	3.68	4.88	0.22	1.65	1.44	0.06	0.23	**		-
NT2RP4000528	9.85	11.61	11.68	7.98	5.49	4.94	7.65	1.72	3.24	*	*	-
NT2RP4000537	22.62	28.04	23.49	9.39	8.19	12.07	6.5	9.43	12.67	**	**	-
NT2RP4000541	6.77	5.01	14.31	6.57	5.43	3.77	2.53	4.6	6.37			-
NT2RP4000543	7.3	6.98	11.29	8.27	7.56	3.29	6.55	3.68	5.22			-
NT2RP4000545	22.52	30.13	32.57	13.3	16.58	18.62	18.71	21.42	13.98	*	*	-
NT2RP4000546	4.78	4.57	9.21	6.74	1.72	3.53	3.64	0.82	4.41			-
NT2RP4000548	11.61	18.79	20.98	6.99	2.21	3.23	4.79	2.09	3.98	*	*	-
NT2RP4000556	15.54	16.2	17.27	7.73	3.7	5.61	5.6	2.76	4.84	**	**	-
NT2RP4000557	11.07	12.42	10.89	8.61	4	8.57	12.02	7.59	12.85			-
NT2RP4000558	14.08	10.84	18.04	10.75	6.08	12.87	11.5	18.9	20.52			-
NT2RP4000560	6.31	4.01	7.32	5.77	2.78	5.38	5.37	6.37	4.62			-

101 / 157

2641

WT2RP4000588	23.22	25.79	25.38	10.18	10.21	8.47	12.24	8.65	11.87	00	00	-	-
WT2RP4000583	5.85	2.98	7.97	6.46	2.06	2.6	3.62	4.57	3				
WT2RP4000585	9.33	19	28.37	6.6	3.13	6.84	10.5	7.88	7.04				
WT2RP4000588	7.43	10.21	10.68	6.68	2.44	1.85	5.44	2.87	5.6	0	0	-	-
WT2RP4000590	6.55	7.08	6.76	6.12	1.79	2.53	5.82	2.45	2.62	0	0	-	-
WT2RP4000599	40.85	42.1	36.12	25.51	13.35	23.08	24.88	16.83	12.91	0	00	-	-
WT2RP4000603	16.28	11.07	12.79	9.86	5.09	7.82	5.81	11.76	8.02	0		-	-
WT2RP4000607	18.04	30.52	24.32	19.53	17.65	25.01	23.84	28.13	28.14				
WT2RP4000614	10.28	8.25	15.91	8.83	8.89	11.83	10.16	8.95	16.38				
WT2RP4000634	6.38	4.3	5.27	7.5	2.83	3.42	3.64	2.83	2.88	0	0	-	-
WT2RP4000638	23.34	31.48	44.47	17.42	11.53	10.82	16.58	9.75	16.96	0	0	-	-
WT2RP4000648	20.24	25.67	26.97	19.17	9.45	12.88	31.67	26.66	31.95	0		-	-
WT2RP4000657	29.17	33.57	37.98	49.48	41.57	42.06	36.04	23.15	32.67	0	+	-	-
WT2RP4000691	5.58	4.18	3.63	5.96	2.37	3.18	5.01	2.34	4.68				
WT2RP4000697	6.38	8.16	10.53	8.34	4.77	12.75	6.28	6.22	8.11				
WT2RP4000704	1.96	1.59	2.11	5.91	1.47	3.12	1.95	2.78	2.48				
WT2RP4000710	20.97	14	15.55	13.68	8.51	18.99	7.53	8.31	11.22	0		-	-
WT2RP4000713	4.84	3.28	2.49	5.57	2.13	4.31	3.32	1.62	2.08				
WT2RP4000724	3.62	3.63	2.74	6.31	1.65	2.66	3.33	4.03	5.81				
WT2RP4000725	3.89	2.42	3.21	6.35	1.7	3.24	4.75	1.9	3.79				
WT2RP4000726	11.58	10.9	20.87	12.34	10.22	7.45	18.8	22.52	15.47				
WT2RP4000737	12.75	11.71	18.21	12.21	6.25	7.06	13.69	12.39	10.01				
WT2RP4000739	8.74	4.93	10.91	9.95	3.78	6.83	6.48	6.27	7.1				
WT2RP4000749	16.48	11.21	22.1	13.84	13.31	13.35	10.75	11.02	11.19				
WT2RP4000759	9.99	10.26	16.79	12.18	3.6	7.13	3.73	6.86	6.05				
WT2RP4000774	8.28	8.05	14.72	7.64	2.48	6.01	4.16	4.89	5.59				
WT2RP4000781	4.61	3.97	6.84	6.54	2.43	3.37	3.39	2.48	3.03				
WT2RP4000783	17.33	30.73	35.48	25.7	12.51	13.28	15.68	12.88	21.97				
WT2RP4000788	24.76	10.56	23.27	28.79	17.17	17.48	11.21	13.65	10.18				
WT2RP4000782	5.04	4.94	10.96	8.67	1.88	4.24	2.65	4.16	5.52				
WT2RP4000809	342.08	122.2	157.03	125.01	165.45	173.38	118.64	164.8	122.66				
WT2RP4000817	8.94	6.91	14.48	7.71	3.01	5.35	5.37	3.83	4.85				
WT2RP4000821	138.37	90.35	131.65	91.89	114.89	72.17	100.95	73.86	85.69				
WT2RP4000822	14.55	12.28	20.9	11.12	8.44	15.29	16.17	13.27	18.76				
WT2RP4000823	2939.8	2567.2	3305.2	1530.8	2690.7	2293.4	1708.4	1550.9	2582.9				
WT2RP4000831	78.61	54.25	80.68	72.05	94.12	99.31	70.84	58.37	82.02				
WT2RP4000833	27.77	14.49	23.94	22.18	7.07	21.11	19.12	24.88	29.44				
WT2RP4000837	6.72	4.87	9.85	8.02	2.92	3.88	2.67	4.16	7.79				
WT2RP4000839	145.23	125.82	232.83	173.61	189.29	177.68	124.91	117.98	102.54				
WT2RP4000846	29.27	18.32	20.88	21.05	17.61	12.28	9.05	14.78	13	0		-	-
WT2RP4000848	18.2	11.65	15.75	19.15	9.44	13.53	19	14.72	12.39				
WT2RP4000855	6.13	7.14	11.29	10.68	4.62	6.82	4.65	3.21	3.53				
WT2RP4000863	6.72	6.6	6.48	8.17	4.81	5.97	4.42	5.29	3.77	00		-	-
WT2RP4000885	45.32	26.52	38.54	39.71	38.1	49.51	43.48	35.35	41.09				
WT2RP4000873	144.06	68.9	135.34	139.48	79.72	155.83	87.22	97.84	116.06				
WT2RP4000874	10.17	6.05	18.44	7.29	2.79	4.52	8.81	5.9	8.53				
WT2RP4000875	14.13	12.82	21.34	9.26	7.3	7.14	8.99	11.91	13.22	0		-	-
WT2RP4000878	26.06	33.12	27.77	33.84	31.46	17.83	20.58	24.01	24.14	0		-	-
WT2RP4000879	8.38	6.53	9.14	6.87	3.61	3.32	2.85	3.38	3.68	00		-	-
WT2RP4000880	35.09	26.71	47.85	60.74	42.76	41.2	28.03	38.06	33.07				
WT2RP4000881	701.32	715.69	430.85	690.64	840.24	746.75	690.32	324.78	792.55				
WT2RP4000894	21.41	14.41	21.14	15.13	8.85	12.55	11.08	18.34	17.24				
WT2RP4000898	2.97	2.72	11.52	4.95	1.01	2.82	2.18	3.29	4.66				
WT2RP4000899	44.53	41.78	65.07	39.31	37.51	42.09	31.81	35.26	42.89				
WT2RP4000907	4.19	4.02	10.25	6.24	3.94	2.33	1.69	3.11	0.84				
WT2RP4000908	14.45	17.83	18.1	10.04	7.21	5.31	4.68	6.8	4.87	00	00	-	-
WT2RP4000910	216.22	171.01	317.29	253.37	255.44	240.15	168.21	189.84	183.37				
WT2RP4000918	3.56	6.94	4.05	7.07	6.01	6.69	5.32	2.08	2.17				
WT2RP4000925	5.57	5.14	8.28	6.79	4.75	4.89	3.2	5.18	4.02				
WT2RP4000927	3.1	2.26	4.02	4.89	1.62	0.68	2.5	1.38	1.2				
WT2RP4000928	9.4	7.34	14.03	9.58	4.88	8.73	4.09	5.17	5.62				
WT2RP4000929	3.84	3.91	13.46	5.95	2.85	3.04	2.12	4.29	1.89				
WT2RP4000946	6.69	8.82	14.08	8.15	4.03	3.77	3.48	3.97	1.29				
WT2RP4000947	3.38	4.07	10.59	4.84	0.29	0.97	0.38	0.58	0				

102 / 157

2642

2643

NT2RP4000949	10.71	9.2	18.65	8.88	2.65	3.74	2.78	1.39	2.41	0	0	-
NT2RP4000950	5.81	4.35	6.4	5.28	2.77	1.08	2.31	1.19	3.37	0	0	-
NT2RP4000951	46.28	48.79	57.49	28.43	48.02	26.11	20.47	30.62	16.93	00	00	-
NT2RP4000952	11.88	9.47	12.39	11.59	10.69	8.83	6.01	4.92	5.77	0	0	-
NT2RP4000973	21.72	18.11	23.28	13.29	14.47	13.18	9.73	13.42	15.34	0	0	-
NT2RP4000975	12.7	11.46	22.76	9.61	5.19	8.19	5.82	10.95	9.25	0	0	-
NT2RP4000979	15.16	9.34	21.89	15.27	12.98	9.04	9.28	10.97	8.24	0	0	-
NT2RP4000984	8.41	8.08	14.46	8.57	1.97	3.73	2.22	3.31	3.09	0	0	-
NT2RP4000986	8.88	5.41	12.86	8.12	4.31	1.27	1.5	2.5	1.16	0	0	-
NT2RP4000988	17.65	11.88	13.93	10.42	7.43	7.53	9.24	12.28	11.72	0	0	-
NT2RP4000989	4.85	7.43	8.1	6.81	2.52	1.81	2.98	2.09	3.85	0	0	-
NT2RP4000990	6.25	7.92	5.23	9.42	4.63	4.12	5.36	2.51	2.44	0	0	-
NT2RP4000994	8.9	9.21	16.92	10.87	4.94	9.69	11.79	23.69	21.99	0	0	-
NT2RP4000996	77.3	49.17	79.6	45.81	34.69	39.35	32.71	38.95	48.1	0	0	-
NT2RP4000997	122.55	129.24	107.05	94.09	70.8	28.94	48.25	80.17	57.14	00	00	-
NT2RP4001001	12.48	18.44	15.8	14.39	8.88	9.17	8.14	8.28	5.33	0	0	-
NT2RP4001004	5.22	3.76	7.06	6.01	1.06	2.64	1.83	1.39	1.05	0	0	-
NT2RP4001008	13.89	13.25	17.25	8.9	7.81	7.87	7.25	6.38	7.3	00	00	-
NT2RP4001009	16.48	20.86	24.07	12.83	13.33	10.38	13	7.19	7.56	0	0	-
NT2RP4001010	12.07	13.84	9.65	9.47	6.84	7.52	5.8	4.44	3.25	00	00	-
NT2RP4001013	109.49	147.49	80.54	50.6	50.74	62.83	38.95	52.28	51.67	0	0	-
NT2RP4001020	20.54	17.88	29.5	8.59	9.72	9.68	7.27	4.7	8.7	0	0	-
NT2RP4001036	9.27	12.23	13.79	8.52	7.83	4.92	8.03	8.75	4.44	0	0	-
NT2RP4001041	38.4	40.27	32.89	14.35	15.84	11.72	23.48	9.92	16.78	00	00	-
NT2RP4001042	15.67	10.3	17.68	10.18	7.53	8.42	5.86	5.84	7.9	0	0	-
NT2RP4001046	38.69	45.09	54.4	22.18	26.84	18.1	31.95	24.71	14.49	0	0	-
NT2RP4001050	14.02	22.04	12.83	10.97	2.91	3.94	9.37	3.69	3.94	0	0	-
NT2RP4001051	21.08	16.13	23.41	17.4	11.41	18.25	15.88	11.32	18.06	0	0	-
NT2RP4001057	5.02	3.25	6.98	7.18	2.13	3.82	4.33	2.83	3.14	0	0	-
NT2RP4001063	6.13	4.37	7.2	7.29	2.91	4.39	2.87	3.18	3.35	0	0	-
NT2RP4001084	9.3	9.96	15.17	12.02	6.98	5.07	7.19	9.48	7.87	0	0	-
NT2RP4001087	10.58	14.37	13.98	8.93	4.54	4.48	7.26	9.18	8.36	00	00	-
NT2RP4001078	4	5.51	5.8	4.49	1.94	2.06	3.68	2.7	1.94	0	0	-
NT2RP4001079	8.77	14.47	11.87	8.88	3.49	1.88	7.38	2.56	7.44	0	0	-
NT2RP4001080	8.56	8.71	5.38	7.24	5.74	5.88	6.43	5.64	6.68	0	0	-
NT2RP4001086	8.1	13.85	11.2	8.73	5.69	7.65	7.59	1.27	6.64	0	0	-
NT2RP4001095	21.9	10.2	20.98	20.1	8.21	22.42	25.97	28.91	25.72	0	0	-
NT2RP4001098	9.87	7.14	14.73	8.69	4.17	6.42	4.58	7.87	4.16	0	0	-
NT2RP4001100	87.64	83.13	87.53	54.87	47.17	71.93	63.14	104.74	100.88	0	0	-
NT2RP4001105	4.82	5.85	6.18	8.6	3.1	3.82	2.72	2.96	3.35	00	00	-
NT2RP4001110	4.35	3.82	3.95	6.4	1.65	2.39	2.88	1.95	2.46	0	0	-
NT2RP4001115	9.9	15.92	16.54	12.73	4.3	5.05	12.1	7.78	11.14	0	0	-
NT2RP4001117	47.61	50.83	48.66	30.24	18.2	17.87	37.4	37.77	12.69	00	00	-
NT2RP4001122	7.49	13.49	11.17	8.93	3.06	4.64	5.55	4.07	4.81	0	0	-
NT2RP4001123	15.99	10.87	15.28	13.98	11.8	8.64	4.92	11.31	11.27	0	0	-
NT2RP4001126	32.2	22.76	27.43	23.25	12.48	17.12	8.88	11.87	12.23	00	00	-
NT2RP4001127	5.7	5.54	5.37	7.81	2.72	4.68	3.57	2.2	3.61	00	00	-
NT2RP4001138	6.85	3.88	7.12	9.78	5.56	5.09	4.09	4.44	3.91	0	0	-
NT2RP4001143	12.5	11.85	11.43	11.35	7.29	7.76	9.34	8.18	11.45	0	0	-
NT2RP4001148	6.15	6.48	10.09	9.88	3.88	5.87	3.55	2.88	4.94	0	0	-
NT2RP4001149	10.56	12.29	14.82	14.78	6.44	4.57	8.21	8.46	7.23	0	0	-
NT2RP4001150	2.08	2.58	2.8	5.53	2.21	2.02	2.88	2.81	5.29	0	0	-
NT2RP4001159	28.55	18.88	30.75	17.14	15.33	13.87	7.91	12.94	18.83	0	0	-
NT2RP4001162	19.05	7.92	22.71	10.51	6.28	6.36	7.29	6.36	7.28	0	0	-
NT2RP4001170	4.82	3.8	7.97	5.23	1.81	3.87	2.53	3.13	2.09	0	0	-
NT2RP4001174	28.86	24.87	30.19	18.82	12.05	22.33	28.65	26.07	34.84	0	0	-
NT2RP4001175	41.57	44.74	48.16	32.54	33.99	41.64	26.8	24.72	38.38	0	0	-
NT2RP4001176	940.52	839.83	1091.8	860.38	609.6	980.82	721.44	858.51	780.83	0	0	-
NT2RP4001184	63.88	45.01	83.95	80.3	52	64.4	59.44	58.17	76.19	0	0	-
NT2RP4001190	36.14	14.18	33.12	25.99	15.11	24.84	10.23	17.15	15.95	0	0	-
NT2RP4001199	7.47	5.68	10.88	10.65	3.07	4.18	4.98	7.07	8.52	0	0	-
NT2RP4001206	64.73	49.7	68.67	51.07	49.14	54.08	37.12	41.81	43.38	0	0	-
NT2RP4001207	7.32	2.58	10.18	7.77	2.4	3.4	3.22	3.79	1.85	0	0	-
NT2RP4001210	10.03	7.34	12.93	9.04	5.66	7.58	3.96	6.73	5.83	0	0	-

103/157

2644

2645

2646

NT2RP4001213	7.81	5.05	8.73	8.57	3.73	3.8	4.8	3.72	3.88				
NT2RP4001214	3.93	5.31	4.9	6.79	4.2	2.68	8.35	2.95	4.58				
NT2RP4001219	16.88	22.38	19.11	23.96	11.89	15.23	12.2	5.76	8.23	*			
NT2RP4001228	31.68	26.26	28.89	28.26	18.75	24.21	24.62	30.13	28.4				
NT2RP4001235	19.17	16.01	20.65	14.75	7.88	8.31	4.98	7.75	11.8	*	*		
NT2RP4001256	7.87	5.89	15.88	8.52	4.69	3.88	3.22	7.66	5.44				
NT2RP4001257	10.17	7.29	12.81	9.33	4.43	6.16	4.58	7.66	4.3				
NT2RP4001260	9	7.35	9.45	7.52	2.82	3.83	1.87	4.5	2.81				
NT2RP4001261	14.73	14	19.92	14.97	13.67	11.83	31.07	30.85	37.38	**			
NT2RP4001274	46.98	48.98	34.12	11.26	8.82	9.09	7.45	8.68	5.77	**	**		
NT2RP4001276	24.77	28.67	28.03	20.15	26.05	19.84	15.05	8.27	8.86	**	**		
NT2RP4001283	274.25	164.44	266.86	208.49	145.59	283.17	127.22	121.38	88.24	*			
NT2RP4001299	70.57	84.2	82.88	48.39	43.31	37.19	12.59	32.14	14.83	**	**		
NT2RP4001313	1.97	3.19	11.08	5.08	1.72	1.86	1.4	3.15	1.49				
NT2RP4001315	13.31	15.88	15.77	13.24	9.24	10.09	4.21	7.24	8.5	*	**		
NT2RP4001320	57.94	48.23	43.52	51.85	61.21	49.26	36.69	35.49	37.13	*			
NT2RP4001325	218.13	214.83	355	324.05	280.73	208.15	240.45	201.7	209.88				
NT2RP4001330	68.24	37.07	81.11	58.79	61.6	67.7	54.68	37.6	43.21				
NT2RP4001339	6.71	9.24	9.28	9.34	4.49	4.79	2.82	4.96	4.68	*			
NT2RP4001343	117.66	68.55	122.62	134.91	97.85	136.85	58.08	81.43	87.93				
NT2RP4001344	91.3	59.49	108.05	186.25	89.38	122.67	65.91	65.77	78.19				
NT2RP4001345	8.99	7.64	17.74	11.99	4.82	8.81	4.98	8.9	5.71				
NT2RP4001351	18.43	14.61	27.02	22.21	18.47	17.51	14.3	12.05	12.26				
NT2RP4001352	3.32	5.15	8.81	5.72	0.75	1.22	1.09	1.47	0.82	*			
NT2RP4001355	4.79	4.23	4.91	4.97	2.19	1.13	2.63	2	2.13	**			
NT2RP4001367	14.2	18.94	22.86	11.42	11.61	7.99	7.87	8	4.49	*			
NT2RP4001372	4.19	2.77	5.41	7.79	3.55	1.81	2.83	1.82	2.32				
NT2RP4001373	61.47	33.24	43.9	18.18	26.86	13.77	10.57	24.44	16.81	*	*		
NT2RP4001375	4.46	4.59	11.77	7.62	2.36	2.63	2.14	3.81	2.97				
NT2RP4001379	5.44	4.91	12.41	7.76	2.24	3.28	2.93	5.12	3.63				
NT2RP4001381	28.02	25.15	21.15	23.38	16.12	13.95	17.03	26.57	24.23				
NT2RP4001386	29.69	19.44	35.04	29.56	18.78	18.25	18.75	24.44	26.65				
NT2RP4001389	11.24	11.58	9.67	9.26	8.9	5.34	3.69	2.72	3.74	*	**		
NT2RP4001396	5.21	4.67	8.34	8.72	2.25	1.78	3.47	3.22	1.99				
NT2RP4001407	5.68	5.49	6.61	7.81	4.64	1.79	3.65	1.8	5.23				
NT2RP4001409	8.48	7.57	16.74	7.93	2.96	5.53	4.77	8.78	4.45				
NT2RP4001410	76.31	89.86	93.62	25.06	52.86	47.85	42.92	82.08	57.7	*			
NT2RP4001414	30.84	57.5	83.9	31.08	28.37	24.44	13.71	23.74	21.51	*			
NT2RP4001424	11.24	11.55	15.27	11.2	4.99	5.97	6.22	6.82	6.88	**			
NT2RP4001433	18.8	14.14	18.87	12.54	11.03	8.58	10.55	8.89	6.43	*	*		
NT2RP4001438	68.21	52.84	81.07	58.91	48.01	54.6	39.37	44.33	43.61	*			
NT2RP4001442	7.93	5.58	7.34	6.34	2.77	3.09	4.34	2.59	3.24	*			
NT2RP4001447	7.95	6.53	5.53	8.92	3.73	3.72	3.38	1.74	2	**			
NT2RP4001468	28.98	30.25	30.86	13.22	11.9	17.04	5.41	8.17	9.81	**	**		
NT2RP4001467	58.03	68.43	78.02	42.37	52.35	39.87	17.7	35.75	28.38	*	**		
NT2RP4001472	23.78	41.35	30	13.57	13.9	13.93	11.58	12.27	10.38	*	*		
NT2RP4001474	7.87	8.46	14.35	7.2	5.31	5.23	4.32	3.7	3.69	*			
NT2RP4001483	6.68	4.49	11.62	7.08	3.5	3.8	3.02	2.79	3.04				
NT2RP4001488	25.21	54.85	68.99	28.07	36.56	22.06	36.35	46.48	23.9				
NT2RP4001492	10.33	5.85	8.11	7.33	2	4.79	7.35	5.55	5.24				
NT2RP4001498	7.69	4.68	4.14	7.14	3.13	3.73	4.3	3.05	2.12				
NT2RP4001502	209.76	206.24	160.31	118.11	105.88	95.3	44.44	56.95	57.25	*	**		
NT2RP4001503	7.44	8.32	13.64	9.86	7.22	6.9	4.97	6.78	6.81				
NT2RP4001507	13.79	9.83	18.56	10.78	5.48	8.77	10.23	16.21	17.48				
NT2RP4001510	5.75	3.91	8.87	6.33	3.38	4.41	4.72	4.9	4.83				
NT2RP4001516	8.14	6.89	9.41	7.53	2.26	3.8	4.88	3.25	3.25	*			
NT2RP4001520	131.31	173.55	237.05	131.12	128.89	116.61	84.53	82.4	103.88				
NT2RP4001523	33.88	38.07	38.29	18.41	8.08	12.46	21.96	12.77	13.03	**	**		
NT2RP4001524	21.05	28.38	29.26	13.82	9.16	8	8.02	5.85	9.68	**	**		
NT2RP4001529	18.64	17.51	18.7	21.14	18.37	22.82	12.83	17.42	17.09				
NT2RP4001531	12.2	10.44	17.59	14.87	8.44	10.35	9.15	12.78	10.56				
NT2RP4001546	209.55	317.08	217.28	217.42	252.35	233.58	165.75	290.78	247.68				
NT2RP4001547	58.85	72.72	70.29	50.48	41.68	51.65	70.8	75.31	83.14				
NT2RP4001551	5.28	7.23	8.07	6.96	2.03	2.81	2.74	2.79	2.73	**			

104 / 157

2647

2648

NT2RP4001554	5.61	4.66	4.4	5.85	1.41	2.54	5.43	2.64	2.46				
NT2RP4001567	14.43	18.57	12.18	9.17	4.93	4.07	5.01	3.84	5.68	*	**	-	-
NT2RP4001568	20.92	33.87	32.23	19.47	17.85	14.08	25.04	13.01	8.31				
NT2RP4001569	38.32	29.49	40	37.57	26.2	39.13	21.18	30.23	30.47				
NT2RP4001571	15.01	7.71	15.89	14.4	7.44	9.88	7.38	9.18	8.27				
NT2RP4001574	25.53	25.21	25.8	22.48	16.52	21.02	13.78	20.38	15.43	*	*	-	-
NT2RP4001575	5.89	6.11	13.98	11.72	4.86	9.13	5.91	5.72	3.19				
NT2RP4001578	138.32	158.83	178.17	106.73	98.03	117.94	140.07	117.64	162.41	*		-	-
NT2RP4001592	9.9	23.21	17.14	17.18	3.08	11.08	13.53	10.53	12.39				
NT2RP4001593	38.07	48.78	48.75	25.68	17.84	15.12	20.23	20.2	20.68	**	**	-	-
NT2RP4001605	8.89	11.76	9.28	8.38	2.88	4.4	4.38	3.06	5.09	*	**	-	-
NT2RP4001606	18.77	11.14	10.78	12.15	11.21	15.99	4.83	13.12	16.6				
NT2RP4001607	8.88	5.47	8.59	11.52	4.98	6.05	4.83	4.03	5.51				
NT2RP4001610	3.61	1.38	4	7.17	1.24	4.9	2.72	3.85	3.08				
NT2RP4001614	9.32	7.4	5.36	10.94	2.88	8.45	8.1	5.66	6.1				
NT2RP4001623	4.1	3.9	7.53	8.2	4.59	6.08	6.02	5.72	5.27				
NT2RP4001626	25.96	32.39	24.45	30.25	8.64	19.85	8.25	9.34	7.78	**		-	-
NT2RP4001634	5.5	5.13	5.73	8.25	3.86	4.21	6.58	3.87	4.85				
NT2RP4001638	6.66	10.41	10.2	8.73	4.08	6.57	10.07	5.69	6.69				
NT2RP4001644	54.14	31.27	53.8	60.84	32.16	47.2	22.38	29.74	30.92				
NT2RP4001648	19.49	10.73	28.93	21.07	12.28	14.22	11.82	16.28	16.41				
NT2RP4001656	13.27	8.07	17.13	8.91	6.79	8.31	7.24	7.88	8.43				
NT2RP4001666	5.12	4.37	7.8	8.14	1.29	4.07	3.21	5.08	2.86				
NT2RP4001670	3.97	4.22	6.38	10.93	2.63	2.89	3.57	4.05	2.82				
NT2RP4001677	60.48	67.87	89.27	37.88	33.85	40.42	60.13	48.04	66.44	*		-	-
NT2RP4001679	63.4	48.03	74.85	69.08	70.98	72.54	80.35	59.3	58.1				
NT2RP4001695	25.41	48.09	50.45	43.83	20.99	24.93	40.59	28.74	39.81				
NT2RP4001696	10.8	7.02	11.48	8.54	3.39	6.61	2.58	6.02	6.23				
NT2RP4001699	7.34	4.14	12.45	8.78	2.82	5.44	4.48	5.75	8.17				
NT2RP4001717	14.1	8.08	13.01	8.07	5.02	5.29	6.2	4.01	4.24	*		-	-
NT2RP4001718	12.35	7.72	11.2	8.9	5.51	7.17	4.84	7.79	8.21				
NT2RP4001726	3.63	2.84	5.27	5.13	1.69	2.38	2.09	2.38	2.12				
NT2RP4001728	27.43	18.04	36.39	17.49	15.05	17.67	15.15	9	13.24				
NT2RP4001730	17.07	14.23	28.12	18.6	22.34	22.39	15.23	16.15	18.5				
NT2RP4001739	13.08	27.52	18.78	28.89	19.24	18.38	17.13	15.51	14.38				
NT2RP4001741	36.21	33.9	43.78	24.77	17.24	19.75	19.65	43.11	31.48	**		-	-
NT2RP4001753	17.2	14.59	18.48	7.48	4.99	6.73	3.91	7.98	13.47	**		-	-
NT2RP4001780	8.38	7.54	12.21	7.55	2.73	5.75	3.77	6.5	5.81				
NT2RP4001787	449.13	316.48	382.02	579.25	624.67	656.78	481.34	491.96	418.4	**	+	-	-
NT2RP4001788	7.47	3.89	8.52	8.8	3.39	5.62	4.59	3.58	5.07				
NT2RP4001795	63.42	46.97	73.29	43.25	32.42	28.05	20.69	20.32	20.73	*	**	-	-
NT2RP4001803	8.84	7.88	10.54	9.34	4	5.93	4.98	3.55	3.59	**		-	-
NT2RP4001805	5.95	5.69	10.38	8.15	3.79	5.48	9.95	6.17	7.95				
NT2RP4001809	55.79	43.66	54.44	43.11	38.38	44.78	37.5	36.54	53.88				
NT2RP4001817	34.65	23.28	22.77	22.82	15.28	16.99	9.01	18.13	18.73	*		-	-
NT2RP4001822	6.52	3.81	10.26	6.23	2.19	2.7	2.67	9.77	2.8				
NT2RP4001823	3.6	2.46	7.23	4.28	1.65	1.68	0.91	2.7	2.37				
NT2RP4001827	92.63	82.49	90.1	26.32	25.71	23.6	20.01	17.73	25.08	**	**	-	-
NT2RP4001828	61.47	38.75	58.29	48.73	53.93	48.7	40.24	45.94	38.84				
NT2RP4001836	44.8	31.89	48.09	48.58	41.3	52.3	39.14	27.95	48.88				
NT2RP4001838	6.15	7.83	5.55	9.67	5.83	6.27	7.38	14.64	4.59				
NT2RP4001841	120.68	83.99	187.18	139.43	130.99	191.25	82.44	82.4	82.84				
NT2RP4001849	5.44	4.87	11.85	5.66	3.18	2.21	2.03	4.71	2.38				
NT2RP4001881	129.57	85.96	121.65	102.03	95.48	118.7	109.17	140.23	134.58				
NT2RP4001877	47.77	41.26	29.9	28.88	24.84	22.31	17.4	14.34	21.88	*		-	-
NT2RP4001879	28.67	26.1	20.85	19.88	15.94	14.99	7.59	15.66	4.85	*	*	-	-
NT2RP4001889	38.95	24.39	38.19	23.27	20.68	27.73	35.81	41.13	32.21				
NT2RP4001893	8.77	9.6	11.68	7.5	6.4	5.73	3.91	4.94	4.11	*		-	-
NT2RP4001895	20.14	21.68	20.63	23.47	19.99	18.99	15.81	11.88	13.51	**		-	-
NT2RP4001898	54.69	54.01	78.1	65.84	57.05	63.77	28.3	45.74	52.57				
NT2RP4001901	17.19	14.63	24	14.51	10.05	13.25	15.49	18.92	20.72				
NT2RP4001910	121.38	105.48	128.64	129.31	162.14	104.65	69.31	116.85	87.82				
NT2RP4001925	17.15	19.32	26.78	14.34	7.91	7.52	5.99	7.67	6.41	*	**	-	-
NT2RP4001928	13.94	13.08	14.09	14.02	5.92	2.48	4.72	6.01	4.88	**		-	-

105 / 157

2649

2650

NT2RP4001927	22.95	20.64	20.03	14.02	11.73	7.04	7.88	6.56	6.03	*	00	-	-
NT2RP4001931	31.71	30.66	39.63	15.12	11.86	11.1	6.42	8.14	7.84	**	00	-	-
NT2RP4001933	124.92	135.92	129.66	117.23	83.62	79.43	61.4	54.06	63.46	**	00	-	-
NT2RP4001938	21.23	9.61	28.3	17.33	8.99	14.74	12.02	19.38	17.02	*	00	-	-
NT2RP4001942	40.71	48.99	41.11	32.73	26.87	27.19	19.11	35.23	32.36	**	00	-	-
NT2RP4001945	3.45	3.93	9.62	6.09	1.55	2.95	2.15	2.81	1.6	*	00	-	-
NT2RP4001946	10.3	8.04	13.38	10.28	3.79	3.79	3.48	4.08	3.41	*	00	-	-
NT2RP4001947	18.45	15.77	21.11	11.34	9.6	7.92	4.68	3.24	4.84	*	00	-	-
NT2RP4001950	5.04	4.87	6.27	8.87	2.41	1.7	3	2.83	1.3	*	00	-	-
NT2RP4001953	26.47	19.18	22.85	22.48	9.35	15.84	17.83	14.03	17.8	*	00	-	-
NT2RP4001966	6.74	6.66	3.65	6.68	3.2	2.07	1.97	1.5	1.8	*	00	-	-
NT2RP4001970	7.84	6.48	14.19	9.18	3.49	5.37	2.98	4.43	3.21	*	00	-	-
NT2RP4001975	18.38	22.03	28.3	17.31	12	7.09	9.7	16.29	13.08	*	00	-	-
NT2RP4001988	38	38.95	48.12	29.2	31.09	30.9	21.07	21.03	23.16	*	00	-	-
NT2RP4001996	24.73	23.55	24.27	15.38	14.66	5.03	8.32	9.1	9.68	*	00	-	-
NT2RP4002014	11.39	15.95	16.62	10.15	7.55	6.98	8.4	7.61	5.82	*	00	-	-
NT2RP4002018	16.73	14.11	11.76	12.18	9.31	11.09	15.09	12.65	8.98	*	00	-	-
NT2RP4002035	5.82	5.03	4.44	7.72	3.9	4.37	6.69	2.8	2.78	*	00	-	-
NT2RP4002043	18.95	25.2	14.18	17.22	10	14.67	15.39	7.04	10.89	*	00	-	-
NT2RP4002046	24.11	28.29	25.84	19.27	12.78	20.33	11.41	17.55	17.78	*	00	-	-
NT2RP4002052	12	15.65	20.88	9.3	9.08	6.85	4.02	3.61	4.24	*	00	-	-
NT2RP4002056	153.48	150.03	95.32	84.85	34.92	59.97	51.1	38.85	53.88	*	00	-	-
NT2RP4002057	40.68	39.2	39.83	19.31	36.56	26.73	27.83	37.55	52.18	*	00	-	-
NT2RP4002058	17.65	14.46	13.17	10.87	3.28	4.52	5.26	2.89	2.9	*	00	-	-
NT2RP4002064	5.16	7.64	7.19	5.68	1.88	1.84	2.82	1.16	2.09	*	00	-	-
NT2RP4002071	19.67	23.04	16.67	12.81	7.48	6.56	6.07	4.85	9.75	*	00	-	-
NT2RP4002075	4.49	2.91	8.1	6.42	3.43	5.27	3.76	2.11	2.28	*	00	-	-
NT2RP4002078	19.28	11.77	21.52	11.04	6.62	7.49	7.2	13.73	14.2	*	00	-	-
NT2RP4002081	26.93	23.83	24.19	19.25	16.65	23.73	20.7	21.48	20.63	*	00	-	-
NT2RP4002083	12.48	9.47	12.44	9.18	6.34	7.67	7.79	10.15	9.98	*	00	-	-
NT2RP4002099	4.69	4.27	5.69	6.55	1.89	1.68	2.88	2.04	1.79	*	00	-	-
NT2RP4002106	48.22	65.51	111.5	22.9	29.08	18.21	28.91	8.21	28.28	*	00	-	-
NT2RP4002111	21.8	28.93	25.85	13.09	13.73	11.32	21.21	10.01	18.62	**	00	-	-
NT2RP4002112	18.8	22.05	20.04	17.73	8.26	13.72	15.37	6.12	8.09	*	00	-	-
NT2RP4002116	131.42	129.9	145.31	124.89	86.8	130.87	116.59	193.03	130.28	*	00	-	-
NT2RP4002122	9.21	7.05	12.08	10.75	5.68	5.1	3.65	7.4	6.23	*	00	-	-
NT2RP4002126	16.25	10.45	28.23	9.01	12.04	10.44	10.14	17.51	8.7	*	00	-	-
NT2RP4002133	59.92	73.72	76.92	61.59	43.1	58.16	80.78	84.66	88.01	*	00	-	-
NT2RP4002136	12.24	16.73	29.34	16.04	8.23	7.88	20.93	16.43	17.62	*	00	-	-
NT2RP4002139	65.38	168.61	135.79	80.21	28.79	27.09	66.97	65.15	29.84	*	00	-	-
NT2RP4002174	10.15	13.02	11.19	12.5	4.36	7.82	8.88	11.55	11.53	*	00	-	-
NT2RP4002185	23.38	25.82	19.72	13.29	6.3	10.89	10.9	7.33	11.09	**	00	-	-
NT2RP4002186	16.88	7.85	12.65	11.9	7.3	14.74	12.83	14.19	17.64	*	00	-	-
NT2RP4002187	28.59	22.57	36.35	26.74	19.38	12.02	11	23.37	23.58	*	00	-	-
NT2RP4002188	5.08	7.53	8.31	11.68	8.59	7.86	4.55	4.87	6.3	*	00	-	-
NT2RP4002199	5.25	3.74	5.62	11.83	2.92	3.8	2.79	4.05	2.9	*	00	-	-
NT2RP4002206	3.81	1.88	2.83	8.57	1.99	1.73	3.4	4.64	2.09	*	00	-	-
NT2RP4002210	6.45	8.2	5.65	5.42	1.97	2.98	3.71	3.22	3.36	*	00	-	-
NT2RP4002222	6.38	6.14	7.04	6.83	3.57	3.25	6.6	5.65	8.67	*	00	-	-
NT2RP4002241	19.38	17.71	14.08	18.18	11.3	8.8	22.47	17.63	22.91	*	00	-	-
NT2RP4002248	36.62	21.31	27.47	30.48	20.08	28.03	19.35	24.1	27.08	*	00	-	-
NT2RP4002250	1.68	1.78	7.25	3.8	1.3	0.25	0.61	0.65	1.92	*	00	-	-
NT2RP4002252	6.84	5.18	13.48	7.69	2.79	5.42	4.19	2.68	3.8	*	00	-	-
NT2RP4002258	35.43	25.77	32.22	22.21	30.41	16.71	15.9	32.72	23.28	*	00	-	-
NT2RP4002288	22.95	20.57	22.28	19.59	16.95	19.18	16.63	15.69	17.98	*	00	-	-
NT2RP4002290	13.7	13.78	25.98	11.44	11.61	14.66	7.05	8.48	10.18	*	00	-	-
NT2RP4002298	10.23	12.39	16.74	11.9	8.3	6.18	9.68	4.97	3.02	*	00	-	-
NT2RP4002306	15.16	15.97	20.14	18.18	10.48	11.51	13.4	11.08	13.62	*	00	-	-
NT2RP4002308	8.45	5.65	13.02	8.55	4	9.83	3.47	4.84	6.83	*	00	-	-
NT2RP4002336	36.27	19.88	28.94	13.28	11.25	15.48	7.56	12.49	16.8	*	00	-	-
NT2RP4002340	2.84	2.81	9.7	4.27	0.88	0.87	1.22	1.79	3.04	*	00	-	-
NT2RP4002361	9.6	7.94	13.29	10.99	5.46	5.67	5.78	8.11	5.48	*	00	-	-
NT2RP4002367	2.79	2.98	7.18	5.28	1.95	2.56	3.24	1.85	1.33	*	00	-	-
NT2RP4002368	24.08	16.92	33.22	37.33	27.67	13.68	19.74	22.15	15.48	*	00	-	-

106 / 157

2651

2652

NT2RP4002377	39.07	34.91	52.74	47.19	54.21	38.25	54.01	36.88	39.01				
NT2RP4002400	8.23	10.17	7.33	8.65	4.11	5.54	5.29	2.67	3.53				
NT2RP4002420	16.72	9.87	22.95	18.38	10.72	16.61	14.74	22.3	29.66				
NT2RP4002432	134.52	97.55	182.62	149.32	142.05	170.39	110.43	102.21	77.71				
NT2RP4002447	15.99	16.51	16.81	15.14	7.65	8.09	7.58	13.4	16.29				
NT2RP4002451	15.11	13.48	18.12	12.6	10.1	7.99	7.15	14.07	11.51				
NT2RP4002481	94.61	109.24	116.21	78.57	91.21	74.75	80.34	65.93	74.08	*	*	-	-
NT2RP4002486	16.89	12.32	18.5	14.11	11.37	5.17	8.32	4.06	4.53	**	*	-	-
NT2RP4002517	13.35	14.55	13.22	12.72	9.03	11.46	10.8	6.18	8.61	*		-	-
NT2RP4002556	14.42	18.7	14.04	18.8	13.59	13.31	18.09	6.06	5.52				
NT2RP4002569	16.26	11.41	27.18	13.39	6.88	10.11	3.13	6.78	11.58				
NT2RP4002587	4.28	5.8	13.49	5.9	21.35	3.55	2.37	3.48	4.03				
NT2RP4002591	11.21	8.73	18.61	10.63	3.25	5.39	2.62	6.88	4.42	*		-	-
NT2RP4002607	7.19	4.54	8.62	7.82	4.54	6.78	3.93	3.64	3.85				
NT2RP4002627	129.04	84.31	131.91	123.08	121.1	113.29	100.69	103.33	117.69				
NT2RP4002628	23.42	19.42	25.54	15.3	10.07	9.36	10.28	14.93	6.74	*	*	-	-
NT2RP4002630	34.52	45.14	38.61	24.65	19.47	21.97	38.2	24.12	21.41	**		-	-
NT2RP4002639	89.58	58.92	77.5	92.97	83.99	82.67	76.62	44.55	74.99	*	+		
NT2RP4002641	10.87	7.59	20.83	9.6	5.64	6.42	5.77	9.03	8.9				
NT2RP4002654	167.82	202.76	171.29	210.71	275.89	155.29	112.71	313.1	177.93				
NT2RP4002668	11.89	14.59	18.31	13.81	8.64	6.37	5.6	13.8	11.32				
NT2RP4002677	38.73	39.84	49.87	21.84	19.97	14	8.4	12.92	7.78	**	**	-	-
NT2RP4002715	50.14	23.66	88.65	78.28	49.48	49.64	41.22	48.79	51.15				
NT2RP4002750	5.11	4.54	6.02	5.02	2.86	2.29	1.61	2.6	1.93	**		-	-
NT2RP4002784	13.9	15.39	21.97	18.43	16.29	10.65	10.61	11.54	8.19				
NT2RP4002791	19.14	18.24	15.43	15.81	11.98	5.78	6.28	6.12	6.25	**		-	-
NT2RP4002811	7.07	4.03	15.62	8.02	4.02	6.88	1.73	4.87	5.66				
NT2RP4002830	22.24	23	26.1	14.14	12.81	9.89	5.48	12.65	7.49	**	**	-	-
NT2RP4002832	6.62	7.2	10.83	9.33	4.27	5.64	7.23	6.78	8.45				
NT2RP4002850	120.6	112.69	112.61	87.95	73.32	75.67	95.07	111.13	108.7	**		-	-
NT2RP4002874	8.25	8.24	13.89	8.67	2.89	3.64	1.21	3.45	2.85	*		-	-
NT2RP4002884	89.55	83.99	93.99	87.7	70.28	83.78	83.11	54.66	68.06	**		-	-
NT2RP4002888	6.66	8.16	10.03	7.45	4.69	3.72	4.88	3.64	2.84	*		-	-
NT2RP4002891	19	20.18	21.15	16.8	8.39	11.81	12.08	8.09	11.53	*	**	-	-
NT2RP4002894	39.88	38.92	34.99	21.18	29.78	22.86	11.88	16.42	39.6	*		-	-
NT2RP4002896	7.81	11.66	17.8	10.13	3.4	6.5	5.01	4.37	5.08				
NT2RP4002905	5.89	3.63	14.35	7.98	3.09	2.88	3	5.21	1.93				
NT2RP4002907	14.62	12.9	19.78	10.14	7.04	6.74	11.88	12.75	9.47	*		-	-
NT2RP5003459	437.62	468.29	451.9	449.8	544.78	414.48	488.17	464	478.98				
NT2RP5003481	21.93	17.54	18.95	12	9.18	10.65	7.44	8.63	4.58	**	**	-	-
NT2RP5003471	62.72	62.97	42.21	31.11	21.1	25.12	48.83	20.37	35.04	*		-	-
NT2RP5003477	5.7	3.4	3.74	5.93	3.44	2.88	3.17	2.23	2.28				
NT2RP5003487	1749.6	1957.3	1778.8	1241.4	1573.1	1807.7	1562.4	2343.9	2243.6				
NT2RP5003492	6.08	6.5	15.88	6.67	3.98	3.86	2.59	3.25	1.92				
NT2RP5003500	4.28	2.52	9.75	8.65	2.34	2.66	2.29	2.63	3.57				
NT2RP5003506	10.49	9.24	16.38	12.74	9.52	9.22	9.69	8.29	9.27				
NT2RP5003512	6.99	7.08	8.63	5.95	2.46	3.38	2.78	2.88	3.49	*	**	-	-
NT2RP5003522	10.07	9.29	8.65	5.18	3.12	3	5.35	1.85	3.07	**	**	-	-
NT2RP5003524	8.44	9.05	5.63	7.13	1.41	1.72	4.68	2.23	3	*		-	-
NT2RP5003527	117.46	141.32	115.3	136.41	95.19	117.09	134.81	57.44	67				
NT2RP5003531	24.59	24.67	34.31	22.85	17.67	26.56	12.06	25.82	35.99				
NT2RP5003534	6.96	7.88	13.1	6.91	5.38	2.92	2.47	4.02	3.38	*		-	-
NT2RP5003539	19.43	27.72	47.01	19.42	30.65	12.48	29.24	22.48	16.27				
NT2RP5003542	12.58	7.81	13.25	8.64	4.5	5.83	7.1	5.6	4.05	*		-	-
NT2RP5003550	5.27	8.68	7.67	5.81	3.59	4.26	5.19	3.06	5.18				
NT2RP5003553	5.9	4.93	5.66	6.03	1.07	1.6	2.58	2.13	3.15	**		-	-
NT2RP5003574	10.12	8.88	7.2	8.89	3.35	3.32	4.78	3.03	4.98	*		-	-
NT2RP5003583	38.48	60.53	36.08	34.94	19.9	25.64	29	16.57	38.99				
NT2RP5003590	9.33	8.85	11.79	8.7	3.93	5.47	3.33	5.64	6.88	*		-	-
NT2RP5003593	7.59	5.31	8.35	10.58	5.68	4.44	3.17	5.91	3.98				
NT2RP5003599	4.7	6.51	7.04	5.86	2.6	3.34	4.3	7.17	3.53				
NT2RP5003601	23.63	19.21	18.37	28.9	18.17	21.68	18.73	17.92	16.85				
NT2RP5003603	8.02	6.6	8.17	6.66	2.33	2.81	2.78	2.24	3.12	**	**	-	-
NT2RP5003608	31.62	37.36	38.66	11.76	5.01	6.27	13.35	6.28	18.64	**	**	-	-

107 / 157

2653.

2654

NT2RP6000182	11.67	12.88	10.81	11.27	4.67	4.18	7.83	8.13	8.65	00	-
OVARC1000001	10.18	10.88	7.82	10.48	3.67	5.27	8.05	6.3	4.32	0	+
OVARC1000003	19.46	16.04	21.53	24.58	18.39	28.35	24.31	27.93	25.27	0	+
OVARC1000004	164.65	181.68	111.37	257.56	204.62	289.69	148.7	142.43	163.8	0	+
OVARC1000006	8.96	9.17	13.23	12.38	8.89	7.89	8.48	11.04	16.91	0	-
OVARC1000011	12.84	15.42	21.83	13.6	7.52	11.75	8.01	7.31	7.45	0	-
OVARC1000014	9.51	5.51	9.75	8.87	3	2.67	4.2	4.5	5.23	0	-
OVARC1000017	7.8	10.89	11.1	9.23	5.72	5.56	7.08	8.83	7.58	0	-
OVARC1000020	39.97	88.58	58.17	35.12	21.04	21.52	23	21.16	19.52	0	-
OVARC1000035	53.62	63.74	58.85	49.18	30.89	52.6	44.01	25.83	35.4	0	-
OVARC1000037	44.18	27.1	38.64	21.27	17.7	38.96	22.76	28.28	23.58	0	-
OVARC1000050	21.2	16.03	18.83	19.43	6.93	18.25	18.85	19.08	20.93	0	-
OVARC1000060	8.69	7.53	9.42	11.74	4.69	15.1	7.88	12.81	10.28	0	-
OVARC1000068	11.63	6.29	9.08	11.58	8.06	9.18	5.71	8.97	7.36	0	-
OVARC1000069	130.28	91.82	110.55	138.44	48.02	81.89	34.87	114.87	91.52	0	-
OVARC1000071	4.7	2.69	6.56	6.14	1.28	3.78	7.5	3.79	3.09	0	-
OVARC1000075	608.15	715.16	979.87	596.89	491.8	521.41	1505.9	1295.8	1454	00	+
OVARC1000083	31.41	22.53	32.47	28.89	18.2	12.73	23.7	17.6	16.65	0	-
OVARC1000085	150.64	132.35	115.14	65.86	79.36	104.26	57.87	55.95	48.39	0	-
OVARC1000086	21.41	22.84	29.16	11.19	15.67	9.71	14.87	19.23	16.21	0	-
OVARC1000087	5.76	4.1	5.37	4.05	5.48	2.89	2.78	3.2	3.4	0	-
OVARC1000090	32.43	34.99	39.46	22.05	34.29	19.79	30.39	42.7	35.58	0	-
OVARC1000091	19.21	16.58	18.4	9.38	12.78	8.63	3.97	5.82	3.09	00	-
OVARC1000092	12.38	12.28	20.09	9.9	9.89	8.81	11.14	12.55	9.97	00	-
OVARC1000105	143.72	148.63	164.07	80.33	78.71	97.82	141.54	139.38	177.6	00	-
OVARC1000106	68.13	60.31	43.64	33.71	26.64	28.64	25.97	22.43	26.91	00	-
OVARC1000109	41.67	37.23	42.5	18.44	22.08	21.68	20.05	26.41	16.1	00	-
OVARC1000113	24.72	16.19	25.29	12.07	22.86	8.61	9.56	13.5	7.9	0	-
OVARC1000114	20.78	17.23	18.89	9.6	15.6	9.38	19.18	20.46	17.77	0	-
OVARC1000133	17.57	29.71	58.68	7.88	21.62	13.9	11.08	7.62	51.45	0	-
OVARC1000137	28.29	21.24	32.71	19.41	23.84	21.06	17.64	15.76	21.29	0	-
OVARC1000139	159.47	147.34	170.25	146.09	161.48	121.94	114.2	88.17	91.75	00	-
OVARC1000145	5.48	5.11	5.8	4.05	7.97	2.4	2.98	4.01	2.81	00	-
OVARC1000148	28.75	26.35	26.82	10.67	7.97	11.29	11.04	13.8	10.7	00	-
OVARC1000151	10.66	7.69	7.49	4.61	6.78	7.44	5.23	8.46	5.82	0	-
OVARC1000157	131.87	72.71	169.18	28.64	46	29.09	32.55	30.3	35.03	0	-
OVARC1000182	5.48	6.74	10.66	5.09	8.45	3.6	3.17	7.57	4.35	0	-
OVARC1000188	20.5	23.24	26.19	11.56	15.79	12.79	11.3	16.45	13.91	00	-
OVARC1000189	67.7	89.5	89.81	59.55	32.42	51.87	35.88	25.97	33.69	00	-
OVARC1000178	7.01	7.74	9.93	3.35	8.18	5.02	5.87	3.4	4.49	0	-
OVARC1000182	6.26	2.63	4.45	2.72	4.87	3.13	2.7	3.24	3.81	0	-
OVARC1000185	14.26	11.19	12.68	5.9	8.25	3.97	5.99	5.58	7.72	0	-
OVARC1000188	19.42	11.08	15.15	6.28	7.14	4.44	6.86	6.47	9.02	0	-
OVARC1000191	10.3	6.61	7.55	5.45	7.49	3.47	4.08	6.65	4.21	0	-
OVARC1000198	22.01	20.32	21.61	13.28	15.24	8.22	13.24	17.04	19.28	0	-
OVARC1000208	131.85	125.05	141.2	188.48	130.32	109.42	107.54	141.64	112.43	0	-
OVARC1000209	37.5	42.42	48.9	35.76	48.16	33.79	26.69	28.98	20.75	00	-
OVARC1000212	14.17	18.08	15.84	10.34	14.11	8.11	6.32	5.07	5.47	00	-
OVARC1000216	7.5	5.29	9.63	2.31	4.11	1.95	4.22	3.32	1.67	0	-
OVARC1000240	38.72	25.44	20.48	13.87	16.01	11.11	15.84	21.45	17.35	0	-
OVARC1000241	9.83	9.47	8.6	4.52	5.75	4.08	4.87	6.55	5.44	00	-
OVARC1000249	12.5	10.38	14.17	6.89	8.18	4.94	7.53	30.07	10.99	0	-
OVARC1000254	211.04	164.06	171.72	162.84	218.18	173	104.59	109.93	114.49	00	-
OVARC1000255	8.62	8.5	15.01	2.8	3.3	2.22	1.58	8.88	2.88	0	-
OVARC1000267	41.48	57.58	54.05	22.69	31.39	23.55	19.25	29.77	21.42	0	-
OVARC1000275	8.58	9.72	13.14	1.2	3.21	1.15	3.78	4.58	2.47	0	-
OVARC1000287	20.12	19.59	16.09	8.41	13.5	9.21	9.7	16.88	9.51	00	-
OVARC1000288	34.41	32.21	34.44	10.83	20.1	12.59	9.34	11.8	18.85	00	-
OVARC1000298	37.69	35.05	40.4	26.93	41.61	28.19	18.03	19.69	20.76	00	-
OVARC1000302	10.27	8.52	8.63	3.64	6.33	2.17	4.38	8.26	5.64	0	-
OVARC1000334	13.62	8.67	11.83	6.89	9.55	6.81	7.04	9.75	5.89	0	-
OVARC1000307	9.2	9.65	13.33	4.89	7.9	7.07	7.34	4.85	4.2	0	-
OVARC1000309	10.77	7.81	10.14	3.65	4.52	3.3	3.45	4.14	3.51	00	-
OVARC1000312	38.52	63.38	48.95	16.29	33.99	22.18	11.44	23.2	11.33	0	-

108 / 157

2655

2656

OVARC1000313	31.08	25.28	22.97	16.85	33.34	18.32	7.95	14.64	18.92	*	-
OVARC1000321	95.57	70.17	108.42	49.21	78.85	38.91	71.72	84.76	66.41	*	-
OVARC1000326	8.36	7.55	9.19	5.89	4.73	3.83	2.8	4.95	2.81	**	-
OVARC1000327	9.84	8.54	7.7	4.58	5.69	4.67	3.24	3.34	3.39	*	-
OVARC1000331	31.51	26.56	17.69	20.2	32.18	25.32	7.77	12.73	9.94	*	-
OVARC1000336	10.5	10.92	9.88	3.58	8.12	5.55	3.82	2.51	3.12	**	-
OVARC1000347	22.28	19.59	19.77	14.38	17.51	15.31	11.64	10.84	10.7	*	-
OVARC1000348	9.03	9.68	11.58	3.27	3.92	2.72	2.5	3.43	2.61	**	-
OVARC1000353	17.87	18.03	15.89	8.09	14.09	9.86	7.93	4.93	7.38	*	-
OVARC1000377	9.05	5.89	9.25	3.23	8.62	3.71	2.88	2.47	4.83	*	-
OVARC1000382	11.38	10.29	10.44	4.87	5.92	3.1	5	4.78	5.93	**	-
OVARC1000384	15.04	15.77	12.95	3.9	4.19	3.88	3.11	4.29	3.9	**	-
OVARC1000401	12.88	8.81	11.8	7.38	8.78	6.88	5.25	4.94	4.67	**	-
OVARC1000406	452.57	398.91	414.58	242.32	241.83	222.43	202.53	242.65	184.03	**	-
OVARC1000407	11.18	13.52	12.76	4.82	6.15	3.43	3.77	3.81	3.74	**	-
OVARC1000408	143.26	162.71	130.24	110.8	163.58	95.03	87.98	95.98	86.83	**	-
OVARC1000410	12.11	11.07	15.74	6.71	11.27	6.87	6.11	5.8	3.42	**	-
OVARC1000411	11.28	8.14	7.71	4.73	4.82	3.45	4.09	4.41	4.58	*	-
OVARC1000414	15.72	17.28	11.48	7.59	8.9	10.02	14.38	14.47	14.88	*	-
OVARC1000420	11.3	11.1	10.67	5.43	7.49	5.25	8.15	6.28	8.64	**	-
OVARC1000421	21.51	18.12	14.88	8.48	10.5	6.1	8.22	6.83	7.06	*	-
OVARC1000427	255.84	237.68	229.09	230.15	285.16	201.09	155.75	128.13	139.78	**	-
OVARC1000431	88.97	107.78	104.05	53.83	78.2	48.18	31.1	18.06	18.58	*	-
OVARC1000437	18.7	18.85	18.29	10.2	15.26	9.01	9.86	5.69	8.45	*	-
OVARC1000438	14.55	21.25	13.48	9.92	17.37	9.21	5.88	6.88	4.08	*	-
OVARC1000440	37.83	35.45	33.24	14.99	19.86	14.94	10.14	8.93	10.09	**	-
OVARC1000442	47.65	39.77	34.85	17.92	25.44	19.14	35.52	28.4	37.91	*	-
OVARC1000443	9.83	12.52	9.02	8.56	33.14	5.85	18.38	5.13	9.47	*	-
OVARC1000451	8.43	5.22	4.46	4.04	5.64	3.64	3.41	4.7	3.49	*	-
OVARC1000455	13.05	12.4	8.82	5.18	5.82	3.89	4.59	3.18	2.41	*	-
OVARC1000458	22.78	19.38	18.72	10.88	17.88	12.45	11.65	7.24	8.12	**	-
OVARC1000467	6.03	5.77	8.99	2.45	3.8	2.2	3.29	2.29	3.38	*	-
OVARC1000470	15.98	13.65	10.41	7.95	9.91	7.26	9.53	10.16	10.25	*	-
OVARC1000473	12.49	10.05	10.69	6.2	7.49	6.37	10.6	7.2	9.15	**	-
OVARC1000479	24.58	30.78	25.75	8.77	13.75	7.98	8.07	9.81	9.07	**	-
OVARC1000484	41.23	29.61	25.62	15.32	17.14	13.32	29.7	25.23	34.72	*	-
OVARC1000485	15.8	17.22	17.4	11.05	13.85	9.12	14.13	11.53	12.18	*	-
OVARC1000496	8.74	8.5	5.88	2.98	8.22	5.46	2.84	3.99	3.9	*	-
OVARC1000520	8.86	5.17	6.53	2.16	5.01	3.05	4.06	3.97	5.92	*	-
OVARC1000522	35.51	29.95	29.3	26.12	30.21	22.71	22.42	16.55	18.08	**	-
OVARC1000526	25.75	25.79	19.89	12.38	14.14	8.5	32.78	19.39	27.42	**	-
OVARC1000529	17.7	16.61	13.49	7.72	9.28	6.61	12.03	7.5	12.38	**	-
OVARC1000533	31.08	27.41	22.68	25.25	24.54	12.97	18.02	13.21	18.14	*	-
OVARC1000543	11.04	5.87	5.45	4.07	4.48	2.25	3.68	3.22	4.84	*	-
OVARC1000550	9.45	7.02	5.98	3.29	4.48	2.68	4.35	2.47	4.43	*	-
OVARC1000553	34.95	36.44	34.04	18.33	24.77	21.97	31.22	31.95	33.2	**	-
OVARC1000556	22.37	23.81	26.15	14.72	11.54	9.38	19.99	16.63	22.58	**	-
OVARC1000557	11.61	9.78	7.49	2.51	3.89	2.24	6.45	4.03	5.39	**	-
OVARC1000581	67.36	70.57	52.24	30.14	35.45	25.39	87.82	47.67	77.14	**	-
OVARC1000584	79.74	89.59	94.8	35.3	37.73	54.01	49.59	52.84	37.28	**	-
OVARC1000573	11.41	12.83	14.29	8.97	10.13	6.51	5.93	10.17	8.55	*	-
OVARC1000576	235.06	210.51	199.48	203.8	268.47	200.45	142.28	126.19	115.56	**	-
OVARC1000578	16.29	13.44	14.45	8.15	12.93	8.58	9.81	12.78	11.9	*	-
OVARC1000581	5.48	3.63	4.65	3.12	3.25	1.85	1.84	1.64	2.35	**	-
OVARC1000585	99.11	60.08	95.7	58.88	53.54	48.9	78.17	82.33	82.01	**	-
OVARC1000588	5.04	6.18	5.9	2.73	4.3	3.79	4.98	4.69	3.2	*	-
OVARC1000605	11.72	9.54	8.98	6.4	6.83	6.02	4.7	4.72	5.55	*	-
OVARC1000622	86.95	79.98	76.79	29.58	85.79	40.9	71.63	83.74	75.18	*	-
OVARC1000636	9.18	7.82	11.99	5.58	6.68	4.67	5.37	6.74	3.7	*	-
OVARC1000640	9.32	12.02	12.8	8.83	10.31	8.57	8.08	8.89	7.4	*	-
OVARC1000649	137.37	159.82	123.85	116.33	170.6	128.38	91.38	85.12	81.65	**	-
OVARC1000651	14.69	15.01	20.85	11.94	15.89	11.61	9.31	9.27	13.9	*	-
OVARC1000677	42.34	38.62	28.23	30.67	34.02	28.61	19.25	22.35	28.93	*	-
OVARC1000678	18.85	21.84	18.64	9.55	15.51	8.53	9.3	11.98	8.4	*	-

109 / 157

【0933】

【表462】

2657

2658

OVARC1000870	17.17	15.32	16.17	8.84	11.11	9.5	8.07	11.07	12.27	**	*	-	-
OVARC1000881	8.2	7.24	6.7	4.69	3.61	5.03	3.58	6.1	3.97	**	*	-	-
OVARC1000882	113.12	97.63	93.09	104.53	135.81	99.85	69.24	48.57	73.95	*	-	-	-
OVARC1000889	57.01	42.21	46.96	44.44	54.09	47.18	30.34	27.71	26.26	*	-	-	-
OVARC1000700	15.99	15.69	20.71	10.16	14.93	8.67	10.07	14.65	13.84	*	-	-	-
OVARC1000703	28.78	29.43	35.22	14.83	20.62	12.69	19.45	19.63	21.91	*	*	-	-
OVARC1000722	101.86	85.68	110.52	89.03	99.63	84.39	80.99	68.44	79.26	*	-	-	-
OVARC1000728	14.33	9.39	12.64	8.8	8.89	8.83	5.72	6.56	7.04	*	-	-	-
OVARC1000727	10.18	7.76	13.09	5.79	7.58	4.18	6.38	6.81	6.67	*	-	-	-
OVARC1000730	29.15	29.95	31.68	12.76	10.82	12.92	11.85	21.92	19.48	**	*	-	-
OVARC1000741	29.23	29.17	34.55	12.03	37.4	9.12	12.19	21.98	17.49	*	-	-	-
OVARC1000746	7.78	5.19	7.06	4.39	4.32	3.45	2.05	2.62	3.13	*	**	-	-
OVARC1000764	38.05	40.13	44.9	31.76	33.38	24.43	25.99	25.42	30.04	*	**	-	-
OVARC1000769	15.32	19.03	26.3	8.64	8.47	6.17	10.92	8.44	11.01	*	*	-	-
OVARC1000771	12.14	10.38	12.38	5.55	9.59	2.99	5.56	6.76	5.39	*	*	-	-
OVARC1000773	208.54	178.97	168.4	156.38	223.88	197.88	120.99	98.51	115.59	**	-	-	-
OVARC1000775	34.98	28.91	27.77	13.91	18.38	12.88	14.63	18.43	17.02	**	**	-	-
OVARC1000778	16.25	14.26	17.29	7.71	11.77	9.22	9.21	13.13	13.25	*	-	-	-
OVARC1000779	5.51	4.08	7.82	1.14	3.76	1.42	1.72	21.49	16.55	*	-	-	-
OVARC1000781	14.72	11.11	14.65	7.53	10.61	4.88	3.59	7.19	4.58	*	**	-	-
OVARC1000787	18.08	11.69	22.35	5.86	7.21	5.45	8	11.11	9.91	*	-	-	-
OVARC1000789	23.78	29.78	21.78	12.72	20.51	12.99	10.1	11.61	10.12	**	-	-	-
OVARC1000800	37.24	62.19	47.92	16.44	23.62	15.47	18.1	25.22	22.47	**	**	-	-
OVARC1000802	10.69	12.79	19.22	6.91	8.97	4.92	12.67	10.28	7.88	*	-	-	-
OVARC1000810	41.47	36.61	45.9	14.82	21.04	14.45	26.69	32.64	43.34	**	-	-	-
OVARC1000811	19.65	16.32	20.3	9.11	11.03	10.04	5.82	8.78	8.16	**	**	-	-
OVARC1000814	47.61	40.61	47.27	21.78	30.43	20.51	28.06	40.57	35.2	**	-	-	-
OVARC1000816	30.44	38.85	45.82	25.26	35.08	20.13	15.73	10.27	23.79	*	-	-	-
OVARC1000817	7.44	5.99	13.01	2.4	3.28	3.96	1.6	3.36	1.57	*	-	-	-
OVARC1000834	9.2	11.08	10.6	3.33	5.92	2.58	4.12	4.4	2.96	**	**	-	-
OVARC1000846	33.2	36.76	34.02	15.54	20.84	16.18	25.4	21.7	22.13	**	**	-	-
OVARC1000850	10.1	9.46	13.84	3.02	6.63	4.83	5.97	6.78	5.48	*	*	-	-
OVARC1000853	124.13	83.82	117.1	60.7	94.43	66.32	58.07	66.48	64.98	*	-	-	-
OVARC1000862	9.56	7.61	8.86	6.22	4.52	5.06	2.49	2.85	3.78	*	**	-	-
OVARC1000873	22.12	9.86	11.25	3.95	10.2	7.01	4.33	4	5.68	*	-	-	-
OVARC1000875	25.32	18.88	20.49	9.43	11.13	12.13	10.43	9.71	12.19	**	**	-	-
OVARC1000876	14.43	11.76	13.8	4.33	8.64	6.48	3.87	5.38	3.7	**	**	-	-
OVARC1000883	16.65	14.83	22.83	10.71	10.48	10.25	4.07	3.52	3.44	*	-	-	-
OVARC1000885	14.94	16.72	15.2	7.47	13.49	6.08	8.84	6.87	4.31	**	-	-	-
OVARC1000886	13.43	12.39	14.55	7.87	12.43	7.38	15.33	6.54	7.36	*	-	-	-
OVARC1000890	238.12	222.49	244.97	168.11	248.06	166.63	120.4	102.6	108.39	**	-	-	-
OVARC1000891	14.84	13	11.09	4.07	5.38	4.65	3.08	4.38	3.01	**	**	-	-
OVARC1000897	6.62	4.78	5.55	2.64	2.48	2.49	2.14	2.77	2.38	**	**	-	-
OVARC1000912	18.82	12.83	16.34	8.16	8.42	8.22	8.21	8.74	10.52	*	-	-	-
OVARC1000914	14.69	8.15	8.14	3.85	5.28	5.96	4.88	6.3	4.1	*	-	-	-
OVARC1000916	17.63	15.34	14.44	6.42	7.74	4.39	7.85	11.07	8.04	**	**	-	-
OVARC1000918	15.19	18.72	21.08	9.85	20.11	11.22	12.02	12.35	10.4	*	-	-	-
OVARC1000924	11.12	12.88	14.01	5.55	9.51	4.44	3.13	4.61	3.65	*	**	-	-
OVARC1000928	46.27	37.28	34.31	5.2	8.81	6.65	4.93	8.15	5.85	**	**	-	-
OVARC1000936	10.69	10.55	8.33	5.2	9.07	4.74	3.02	5.31	5.15	**	-	-	-
OVARC1000937	7.99	8.04	6.58	4.25	5.19	2.94	3.85	3.95	5.45	*	-	-	-
OVARC1000945	15	14.98	13.39	5.71	5.82	7.78	3.1	4.11	2.76	**	**	-	-
OVARC1000948	7.88	5.08	4.83	2.85	3.28	2.79	2.89	2.39	2.47	*	*	-	-
OVARC1000956	14.68	11.07	13.93	5.16	8.05	6.51	8.38	4.46	6.46	**	*	-	-
OVARC1000958	16.85	16.96	14.18	8.84	10.41	6.42	15.3	9.78	14.01	**	-	-	-
OVARC1000980	47.32	52.37	41.49	22.16	32.89	19.82	46.28	34.15	44.56	*	-	-	-
OVARC1000984	306.84	272.77	304.89	274.81	264.01	211.54	160.52	108.2	102.38	**	-	-	-
OVARC1000971	7.74	8.08	3.76	3.85	7.47	4.63	2.45	4.82	2.32	*	-	-	-
OVARC1000976	109.05	95.15	91.82	89.45	126.14	83.84	64.44	48.09	54.61	**	-	-	-
OVARC1000976	8.41	7.34	6.83	3.4	4.98	2.89	5.48	3.58	2.85	*	-	-	-
OVARC1000981	24.82	24.19	24.61	14.13	15.11	11.79	18.79	11.54	16.77	**	*	-	-
OVARC1000982	21.37	12.18	12.3	7.88	18.73	9.88	5.87	4.64	5.45	*	-	-	-
OVARC1000984	10.44	9.85	10.17	2.64	4.79	2.24	2.88	2.32	3.54	**	**	-	-
OVARC1000995	27.67	24.28	19.39	14.42	17.35	10.4	23.31	17.65	22.26	*	-	-	-

110 / 157

2659

2660

OVARC1000996	12.75	15.42	13.22	7.58	7.82	4.88	10.5	9.45	10.8	**	*	-	-
OVARC1000999	77.92	64.11	52.33	34.81	41.57	26.78	51.13	38.47	53.64	*	-	-	-
OVARC1001000	34.25	25.88	23.55	13.32	17.72	11.08	24.85	23.48	28.11	*	-	-	-
OVARC1001004	4.45	4.47	4.74	3.58	4.7	2.11	3.94	2.09	3.24	*	-	-	-
OVARC1001010	7.8	4.93	6.92	2.94	3.81	2.97	3.34	2.1	2.69	*	*	-	-
OVARC1001011	10.58	5.98	7.28	2.55	4.47	3.01	5.82	4.52	6.68	*	-	-	-
OVARC1001030	135.15	261.75	227.94	119.83	118.81	67.28	114.11	94.32	117.34	**	-	-	-
OVARC1001032	10.53	7.53	10.42	4.8	4.01	3.78	8.13	8.94	7.33	**	-	-	-
OVARC1001034	10.83	11.24	8.27	5.28	4.2	4.56	7.53	5.89	7.35	**	*	-	-
OVARC1001038	35.07	37.32	30.24	14.88	24.71	11.8	14.41	10.88	18.24	*	**	-	-
OVARC1001040	40.95	33.02	38.27	20.4	24.74	15.2	38.11	31.08	34.73	**	-	-	-
OVARC1001041	28.68	24.51	24.41	14.12	19.17	10.91	20.98	17.34	23.83	*	-	-	-
OVARC1001044	8.68	8.02	8.34	3.74	2.78	3.79	6.52	3.05	4.17	**	-	-	-
OVARC1001049	23.67	22.64	22.97	8.18	12.73	9.28	19.48	22.57	21.53	**	-	-	-
OVARC1001051	381.08	484.89	380.08	354.82	548.22	323.29	741.7	583.07	703.22				
OVARC1001054	6.6	5.59	5.17	3.05	5.01	2.19	3.89	3.27	2.79	**	-	-	-
OVARC1001055	14.4	14.88	14.95	7.9	10.31	6.02	5.76	7.88	6.8	**	**	-	-
OVARC1001082	17.05	13.11	19.77	9.47	7.02	5.92	5.47	7.54	3.76	*	**	-	-
OVARC1001085	58.8	47.67	46.94	41.39	58.25	31.45	28.27	24.21	23.74	**	-	-	-
OVARC1001088	12.24	9.85	10	9.05	10.45	10.03	4.48	8.42	6.8	*	-	-	-
OVARC1001072	5.2	5.85	6.88	2.88	7.45	1.74	3.98	4.57	2.54	*	-	-	-
OVARC1001073	7.5	8.58	12.65	4.76	8.4	4.85	5.22	6.98	5.47	*	-	-	-
OVARC1001074	4.7	5.22	8.88	3.02	6.61	3.48	7.1	3.06	2.4	*	-	-	-
OVARC1001076	12.05	11.95	9	4.87	9.88	3.58	7.61	7.71	7.4	*	-	-	-
OVARC1001086	11.18	10.68	13.33	5.34	5.78	6.25	7.82	7.57	6.14	**	*	-	-
OVARC1001088	9.39	8.11	10.75	4.2	10	3.47	6.76	8.44	4.74	*	-	-	-
OVARC1001091	241.22	186.8	193.81	163.03	271.85	178.88	140.22	106.42	112.84	*	-	-	-
OVARC1001092	18.15	15.63	18.59	22.78	28.04	21.17	13.63	15.45	14.87	*	+	-	-
OVARC1001104	8.1	6.77	10.2	3.75	7.44	6.44	3.73	7.33	6.12	*	-	-	-
OVARC1001107	107.49	92.87	112.02	98.9	125.4	90.42	58.37	48.66	58.61	**	-	-	-
OVARC1001113	4.26	4.3	6.18	3.48	7.31	2.59	2.8	2.95	2.48	*	-	-	-
OVARC1001117	14	17.53	15.78	9.22	10.42	7.27	18.8	18.88	12.72	**	-	-	-
OVARC1001118	28.39	25.32	27.08	13.26	13.63	13.64	14.58	18.84	21.18	**	*	-	-
OVARC1001125	15.08	9.24	18.28	5.27	6.72	6.58	3.81	5.47	3.35	*	*	-	-
OVARC1001129	8.66	7.08	12.48	3.09	6.25	3.24	5.49	10.45	6.05	*	-	-	-
OVARC1001132	9.08	13.23	15.1	8.65	15.2	9.07	5.57	13.13	8.52	*	-	-	-
OVARC1001138	155.24	193.79	160.3	115.39	149.38	101.68	122.48	184.58	137.03				
OVARC1001141	12.99	19.8	12.94	7	9.03	7.57	7.48	10.75	7.3	**	*	-	-
OVARC1001154	128.63	107.01	106.33	119.46	158.63	83.66	81.42	78.16	83.39	*	-	-	-
OVARC1001181	17.68	27.02	21.1	11.55	22.82	12.42	16.84	15.27	15.42	*	-	-	-
OVARC1001182	21.74	16.75	19.82	10.25	11.33	9.72	10.1	12.69	11.14	**	**	-	-
OVARC1001183	18.91	8.03	10.73	4.37	5.95	4.38	3.85	7.35	4.62	**	*	-	-
OVARC1001187	38.47	38	43.81	17.29	22.87	18.91	28.74	29.72	35.08	**	*	-	-
OVARC1001189	11.94	7.48	10.4	5.85	13.24	4.72	3.88	5.51	4.65	*	-	-	-
OVARC1001170	28.22	28.34	31.69	14.48	15.02	12.38	12.5	14.3	11.17	**	**	-	-
OVARC1001171	143.81	238.48	129.81	114.4	117.18	158.91	77.44	54.36	128.19				
OVARC1001173	24.91	33.8	35.93	11.93	16.28	13.53	26.95	27.14	25.91	**	-	-	-
OVARC1001176	209.32	244.43	234.51	205.75	340.26	228.89	167.09	178.74	200.74	*	-	-	-
OVARC1001180	67.2	61.17	68.56	20.38	32.42	26.76	42.84	47.13	54.33	**	*	-	-
OVARC1001188	18.84	14.15	17.08	4.37	15.93	5.9	10.16	38.68	8.68	*	-	-	-
OVARC1001200	14.51	7.89	14.82	6.11	7.72	3.65	4.15	15.56	8.11	*	-	-	-
OVARC1001202	37.48	21.56	29.17	10.08	10.43	13.25	17.78	23.14	18.29	*	-	-	-
OVARC1001206	8.83	6.97	11.31	3.08	5.86	2.9	2.03	8.52	4.3	*	-	-	-
OVARC1001209	58.31	51.58	60.2	67.91	69.04	54.41	38.32	28.24	28.87	**	-	-	-
OVARC1001219	14.82	14.11	12.95	3.85	7.48	4.85	4.34	7.98	3.54	**	**	-	-
OVARC1001222	17.11	17.29	18.38	6.81	16.42	9.84	6.58	7.21	5.58	**	-	-	-
OVARC1001232	34.23	34.49	35.17	16.43	23.42	18.3	24.37	27.16	28.45	**	**	-	-
OVARC1001240	22.68	22.79	23.45	11.83	13.49	10.07	14	20.54	14.23	**	*	-	-
OVARC1001243	8.41	5.82	7.88	3.1	7.95	3.99	3.48	18.25	1.41	*	-	-	-
OVARC1001244	48.28	22.42	33.3	20.02	23.25	15.67	15.69	21.46	16.15	*	-	-	-
OVARC1001248	181.8	262.18	216.69	139.78	282.82	181.43	261.49	147.46	197.54				
OVARC1001247	38.18	25.43	28.03	8.87	21.81	17.23	8.81	11	7	**	-	-	-
OVARC1001280	18.58	9.81	13.5	4.35	12.48	3.58	7.06	8.38	13.07	*	-	-	-
OVARC1001281	18.88	9.17	11.3	5.17	7	3.2	4.94	4.18	3.51	*	**	-	-

111 / 157

2661

2662

OYARC1001268	91.7	87.42	103.18	60.09	81.38	43.24	78.88	80.56	58.28	**	-
OYARC1001270	7.04	6.55	7.37	4.89	4.4	3.37	2.38	3.78	2.68	**	**
OYARC1001271	18.59	14.31	18.8	9.15	12.99	7.59	10.4	9.45	13.28	*	*
OYARC1001282	9.08	5.35	8.29	3.47	3.89	2.88	2.79	2.02	1.62	*	*
OYARC1001286	8.37	7.88	7.76	3.73	3.83	3.51	3.2	2.62	1.42	**	**
OYARC1001305	8.13	8.38	10.05	2.51	4.13	1.81	1.91	3.24	2.11	**	**
OYARC1001314	6.35	7.88	10.42	5.15	8.88	5.8	5.8	2.73	3.16	*	*
OYARC1001316	9.21	9.19	7.4	3.28	5.16	2.41	2.09	2	2.8	**	**
OYARC1001329	66.76	60.97	62.75	30.71	38.83	28.85	49.5	54.71	58.08	**	*
OYARC1001380	6.84	7.09	8.03	3.83	3.92	2.72	1.52	3.71	2.82	**	**
OYARC1001335	20.53	13.52	17.21	8.37	10.93	7.02	7.67	13.22	12.68	*	*
OYARC1001338	9.67	3.35	7.39	3.31	2.87	2.49	3.22	2.79	2.03	*	*
OYARC1001339	93.84	87.95	84.08	45.91	68.98	50.22	43.37	46.73	40	**	**
OYARC1001340	8.83	5.17	8.34	3.19	3.73	1.7	2.84	3.21	3.18	*	*
OYARC1001341	30.38	40.21	33.21	15.81	17.58	18.77	23.98	16.89	14.16	**	*
OYARC1001342	319.96	384.64	288.02	528.73	514.44	449.65	353.78	251.2	152.62	*	+
OYARC1001344	24.05	19.69	20.03	8.91	10.1	7.88	18.88	12.81	15.85	**	*
OYARC1001357	8.41	18.75	6.59	4.08	6.58	3.73	3.75	4.14	3.16	*	*
OYARC1001359	58.32	54.08	31.23	23.17	43.75	46.21	51.59	26.81	37.78	*	*
OYARC1001360	5.67	4	3.9	1.82	2.67	1.87	1.89	2.07	1.71	*	**
OYARC1001369	7.8	8.8	8.35	5.24	6.18	2.48	5.5	3.41	1.68	*	*
OYARC1001372	9.87	8.14	5.08	1.49	5.78	2.2	2.9	3.07	0.98	*	*
OYARC1001376	2.27	25.34	21.25	12.73	15.31	10.9	17.23	14.16	15.91	**	*
OYARC1001381	38.84	52.88	35.2	18.78	31.21	17.84	39.83	32.09	35.47	*	*
OYARC1001391	4.35	4.74	4.59	3.41	5.81	3.07	2.98	3.15	1.97	**	*
OYARC1001392	11.01	13.87	11.28	7.11	8.72	6.38	4.32	4.97	3.08	*	**
OYARC1001399	7.11	9.27	5.73	3.82	4.78	3.62	4.04	3.29	4.7	*	*
OYARC1001417	8.85	8.11	7.05	3.89	5.18	3.15	2.59	2.11	3.11	**	**
OYARC1001419	9.55	7.34	7.59	3.89	5.75	3.4	4.52	2.92	1.58	*	**
OYARC1001425	13.5	5.66	6.59	3.04	5.54	5.43	3.86	1.31	1.7	*	*
OYARC1001486	14.91	13.09	11.35	5.12	5.94	3.89	4.58	3.69	3.49	**	**
OYARC1001442	5.57	6.26	6.22	3.11	6.58	2.54	5.62	2.71	2.72	*	*
OYARC1001451	30.68	20.35	20.83	10.67	21.04	10.84	19.07	21.71	23.53	*	*
OYARC1001452	18.95	14.67	11.44	7.52	6.15	5.24	7.7	11.38	6.84	**	*
OYARC1001453	7.88	6.08	6.61	4.22	6.52	4.99	4.72	4.88	4.78	*	*
OYARC1001476	20.63	24.57	30.44	15.92	22.32	7.34	11.05	12.16	18.07	*	*
OYARC1001480	7.35	4.8	7.98	3.71	4.84	5.18	2.87	2.08	3.18	*	*
OYARC1001489	11.26	5.88	6.53	8.18	19.5	4.38	4.69	2.8	2.81	*	*
OYARC1001493	15.28	11.38	13.67	4.88	4.94	2.78	9.11	6.21	8.31	**	*
OYARC1001496	32.05	27.24	23.8	12.75	20.84	11	9.38	5.04	10.85	*	**
OYARC1001499	5.67	6.05	4.7	2.72	3.78	1.85	2.85	4.6	2.67	*	*
OYARC1001505	31.78	25.87	22.22	8.57	18.43	9.44	18.79	15.08	20.69	*	*
OYARC1001508	17.78	13.64	12.09	7.08	8.79	6.78	15.13	14.75	21.14	*	*
OYARC1001510	8.59	5.24	6.08	3.07	5.95	3.22	7.27	7.67	4.6	*	*
OYARC1001516	10.81	12.88	13.12	5.8	9.08	8.84	8.79	7.64	9.81	*	*
OYARC1001525	7.75	5.51	6.34	3.13	4.32	3.68	4.26	3.77	3.9	*	*
OYARC1001542	30.12	29.23	28.39	14.89	25.08	14.49	18.28	13.2	26.28	*	*
OYARC1001544	27.99	28.47	20.43	12.55	18.48	10.77	38.58	24.05	33.39	*	*
OYARC1001548	6.07	8.98	10.22	3.38	5.35	8.68	3.38	3.45	4.28	*	*
OYARC1001547	4.36	6.21	6.19	3.95	3.95	1.63	2.77	2.7	2.24	**	*
OYARC1001555	137.04	110.5	138.25	120.38	174.1	119.31	90.68	69.83	70.87	*	*
OYARC1001580	10.64	9.21	9.05	6.2	10.38	4.19	4.33	6.13	2.34	*	*
OYARC1001589	28.34	25.69	28.31	17.71	18.82	21.58	15.15	17.9	15.82	**	**
OYARC1001570	8.57	12.59	13.54	7.21	9.92	8.14	6.01	6.08	8.45	*	*
OYARC1001577	10.2	13.78	13.09	13.59	15.75	9.11	10.58	8.87	5.25	*	*
OYARC1001578	0	0.27	1.12	0.09	2	0.48	1.37	0.77	0	*	*
OYARC1001595	18.44	38.88	31.49	14.48	23.49	16.88	13.34	13.82	12.27	*	*
OYARC1001800	12.84	12.05	12.79	8.39	9.83	7.52	7.16	10.38	7.31	**	*
OYARC1001607	17.69	15.72	17.84	10.9	18.71	12.05	10.18	16.33	11.6	*	*
OYARC1001610	7.42	7.24	11.17	5.09	7.94	4.34	2.78	4.22	3.65	*	*
OYARC1001811	3.52	5.78	9.33	2.82	5.44	3.47	2.68	4.9	2.78	*	*
OYARC1001615	7.22	4.15	9.71	3.66	4.51	3.88	4.88	4.91	5.78	*	*
OYARC1001836	5.13	4.74	6.28	4.34	6.6	4.17	8.58	8.24	9.12	*	*
OYARC1001854	40.93	55.07	40.85	20.99	33.29	18.88	40.58	43.41	37.28	*	*

112 / 157

2663

2664

QVARC1001702	9.4	5.83	9.29	4.18	2.13	4.1	3.32	3.32	2.46	*	*	-
QVARC1001703	7.15	7.4	12.31	4.47	5.7	4.5	5.65	8.24	3.65	*	*	-
QVARC1001710	18.68	12.34	27.68	8.37	10.17	11.08	8.89	17.12	13.23	*	*	-
QVARC1001711	18.7	15.2	21.81	10.78	18.28	9.63	7.19	14.6	8.6	*	*	-
QVARC1001713	106.12	95.89	110.84	111.33	127.16	96.1	59.14	82.39	67.07	**	**	-
QVARC1001726	7.41	8.41	15.02	8.01	22.48	3.17	4.32	4.47	6.9	*	*	-
QVARC1001726	9.14	11.78	13.18	5.3	10.91	4.63	7.72	5.13	7.08	*	*	-
QVARC1001727	8.13	4.38	5.44	2.42	3.92	3.64	2.37	4.79	3.09	*	*	-
QVARC1001731	328.35	314.92	333.21	240.06	116.88	168.02	275.41	235.38	284.78	*	*	-
QVARC1001735	10.45	7.4	10.3	3.28	8.6	5.18	2.85	50.48	3.58	*	*	-
QVARC1001741	51.89	35.06	44.89	30.3	40.05	30.18	29.82	27.49	31.74	*	*	-
QVARC1001745	48.03	48.42	50.14	20.26	24.89	18.25	35.38	41.14	42.18	**	**	-
QVARC1001759	8.58	9.79	14.25	4.9	9.68	2.64	3.21	5.87	3.68	*	*	-
QVARC1001762	16.44	18.78	16.88	4.89	8.2	5.87	5.45	4.57	3.81	**	**	-
QVARC1001766	83.38	85.91	87.93	59.55	79.17	53.32	53.78	49.12	57.88	**	**	-
QVARC1001767	7.04	5.21	8.42	2.98	5.88	2.78	2.97	1.63	2.74	*	*	-
QVARC1001768	14.52	13.49	18.22	6.31	12.6	7.55	3.97	8.24	6.88	**	**	-
QVARC1001770	28.33	28.28	29.2	18.88	20.54	11.13	6.52	15.1	8.61	*	**	-
QVARC1001776	7.09	4.63	7.55	4.18	5.49	2.76	2.6	3.29	2.71	*	*	-
QVARC1001791	10.39	7.82	10.63	6.37	4.38	3.31	3.58	4.2	5.4	*	**	-
QVARC1001795	7.98	5.23	12.47	1.59	4.12	3.1	3.31	4.76	7.14	*	*	-
QVARC1001798	40.84	44.28	46	20.27	29.31	16.28	30.29	32.18	29.44	**	**	-
QVARC1001802	21.47	24.32	21.73	9.56	17	10.13	12.8	16.34	14.18	*	**	-
QVARC1001825	21.41	25.19	22.93	13.01	23.64	11.53	6.55	6.13	2.98	**	**	-
QVARC1001827	9.03	4.95	8.69	4.35	5.41	3.68	4.23	6.47	5.73	*	*	-
QVARC1001830	149.48	129.88	172.57	133.13	197.68	130.82	61.18	73.6	73.85	**	**	-
QVARC1001812	14.81	11.9	18.98	7.98	7.78	6.77	7.18	8.19	7.68	*	*	-
QVARC1001813	18.16	13.4	12.96	5.79	9.88	8.64	9.17	11.83	7.83	*	*	-
QVARC1001820	13.24	13.12	21.68	6.48	7.84	7.81	9.43	11.92	8.28	*	*	-
QVARC1001828	7.36	8.18	15.13	2.53	4.48	3.18	2.97	2.27	1.93	*	*	-
QVARC1001833	7.55	8	10.4	3.73	5.88	2.72	2.14	3.67	2.55	*	**	-
QVARC1001839	9.51	8.23	9.47	3.48	6.12	2.44	2.33	1.23	3.33	*	**	-
QVARC1001846	10.75	8.47	12.88	4.44	5.49	4.43	2.19	5.18	4.87	*	*	-
QVARC1001849	21.23	22.64	18.18	12.84	14.77	8.26	9.88	17.09	10.17	*	*	-
QVARC1001861	14.15	12.35	14.96	7.88	9.2	8.08	6.33	6.39	8.07	**	**	-
QVARC1001873	10.37	9.13	10.41	3.68	3.81	3.28	3.75	2.72	3.55	**	**	-
QVARC1001879	12.35	13.47	14.5	4.79	6.62	4.85	9.81	8.26	4.82	**	*	-
QVARC1001880	17.98	18.04	17.13	6.7	12.3	8.25	9.7	7.19	8.89	*	**	-
QVARC1001883	10.68	7.64	9.62	5.88	5.31	4.98	4.5	5.37	3.66	*	**	-
QVARC1001900	21.57	23.04	21.79	9.35	13.16	8.54	4.93	5.9	5.98	**	**	-
QVARC1001901	6.1	3.23	6.27	3.78	3.35	2.44	2	2.36	2.55	*	*	-
QVARC1001911	5.85	5.08	5.24	3.87	3.1	3.23	3.29	4.37	2.41	**	*	-
QVARC1001918	15.55	14.04	13.05	10.21	8.28	9.44	6.7	7.73	11.27	**	*	-
QVARC1001928	4.28	4.17	5.98	2.82	2.37	1.14	3.65	1.6	1.43	*	*	-
QVARC1001937	25.6	15.71	13.44	8.48	11.18	11.27	11.49	12.97	7.51	*	*	-
QVARC1001940	8.78	8.78	8.37	2.83	3.28	2.64	2.16	2.98	2.02	**	**	-
QVARC1001942	20.24	15.18	17.14	7.55	9.31	5.48	5.3	6.71	7.34	**	**	-
QVARC1001943	29.63	32.41	20.55	12.87	22.82	16.27	4.34	4.72	5.72	**	**	-
QVARC1001948	14.19	11.75	10.33	3.19	5.85	4.46	2.89	5.47	5.47	**	**	-
QVARC1001950	13.94	15.9	10.94	6.19	8.29	8.75	8.04	7.58	8.81	*	*	-
QVARC1001952	122.9	130.05	103.73	80.58	135.58	102.89	79.11	70.86	72.15	**	**	-
QVARC1001954	7.45	6.81	6.15	3.11	3.63	2.91	2.32	2.33	2.18	**	**	-
QVARC1001983	12.44	11.79	17.81	8.04	6.32	5.28	6.38	9.47	7.04	*	*	-
QVARC1001983	25.71	30.08	25.81	12.15	15.68	10.47	16.02	11.31	11.87	**	**	-
QVARC1001987	13.81	15.75	12.25	8.15	7.8	4.37	5.37	5.81	4.93	**	**	-
QVARC1001989	18.41	18.89	13.98	7.74	12.38	8.71	14.94	11.39	11.29	*	*	-
QVARC1001991	12.31	10.21	10.74	6.47	8.3	7.41	6.69	5.19	5.31	*	**	-
QVARC1002005	33.83	38.39	35.82	17.89	27.08	20.01	28.18	23.73	27.11	**	**	-
QVARC1002048	25.13	24.53	23.58	14.08	17.38	11.27	23.29	17.04	24.48	**	**	-
QVARC1002048	78.05	107.27	88.51	49.63	77.97	65.8	68.65	65.46	67.43	*	*	-
QVARC1002050	9.22	6.62	10.19	4.05	4.76	4.15	4.13	2.81	4.5	*	*	-
QVARC1002058	18.18	18.91	17.55	13.19	18.87	8.99	24.17	22.71	28.4	*	*	+
QVARC1002066	67.02	72.72	47.65	37.79	39.58	35.74	82.17	60.77	51.33	*	*	-
QVARC1002082	34.4	33.12	24.78	17.13	27.63	15.59	48.32	29.05	33.18	*	*	-

113 / 157

2665

2666

QVARC1002011	18.78	18.82	16.41	10.02	4.76	6.56	9.88	8.62	10.94	**	**	-	-
QVARC1002012	11.35	7.73	4.3	3.87	5.5	2.78	3.85	4.58	3.79				
QVARC1002013	11.71	8.5	20.68	4.98	14.44	7.82	5.85	5.4	7.08				
QVARC1002014	8.29	8.49	8.65	2.89	4.04	2.24	4.34	1.07	3.08	**	**	-	-
QVARC1002107	13.48	8.34	12.83	6.08	7.84	8.33	9.41	7.18	12.5	*		-	-
QVARC1002112	28.38	33.01	27.23	12.79	21.35	12.35	12.54	8.82	14.29	*	**	-	-
QVARC1002126	65.18	69.33	51.38	44.58	51.62	38.84	65.71	52.52	60.88	*		-	-
QVARC1002127	6.15	6.37	5.32	1.95	4.41	3.09	5.22	3.48	3.37	*	*	-	-
QVARC1002135	5.22	5.32	3.68	5.18	3.08	2.95	2.58	2.28	2.94	*		-	-
QVARC1002143	7.3	4.6	8.7	3.29	5.14	1.8	3.79	3.84	5.8			-	-
QVARC1002156	14.33	13.88	13.02	10.03	12.28	6.78	8.59	6.29	5.87	**		-	-
QVARC1002158	9.16	5.03	4.85	1.18	3.11	2.41	3.88	2.88	4.37			-	-
QVARC1002165	67.03	61.88	64.25	23.78	38.97	31.97	46.18	44.24	42.15	**	*	-	-
QVARC1002175	10.61	13.89	9.02	5.22	6.29	4.46	9.03	7.11	8.08	*		-	-
QVARC1002178	11.02	11.22	8.01	3.18	5.47	2.82	6.17	3.40	3.57	*		-	-
QVARC1002182	6.32	5.82	5.4	2.92	5.92	3.91	6.46	4.29	5.82			-	-
QVARC1002185	13.32	14.59	15.01	5.81	10.11	6.85	7.78	7.13	8.82	**	**	-	-
PLACE1000064	9.74	9.21	11.81	4.85	7.73	5.01	4.83	8.27	4.83	*	*	-	-
PLACE1000065	27.12	18.01	21.45	13.29	16.54	9.78	14.01	12.75	12.04	*		-	-
PLACE1000066	91.12	67.04	65.23	34.88	62.28	49.96	6.88	11.98	13.8	**		-	-
PLACE1000077	7.28	8.78	8.28	8.37	10.72	8.65	3.04	5.48	2.7	*		-	-
PLACE1000014	2.23	2.04	4.15	1.42	4	3.02	1.88	2.43	1.7			-	-
PLACE1000031	25.49	34.88	32.13	13.87	27.22	11.3	27.49	25.37	24.47			-	-
PLACE1000033	18.11	15.97	17.95	12.01	14.01	16.72	14.29	10.78	15.77			-	-
PLACE1000040	8.82	8.21	10.8	5.78	10.45	6.53	3.81	7.12	4.69	*		-	-
PLACE1000045	20.38	17.3	18.51	15.36	23.83	12.54	8.78	8.57	6.66	**		-	-
PLACE1000050	5.4	7.04	6.5	2.42	4.88	4.46	3.72	6.27	1.7			-	-
PLACE1000051	8.33	8.81	17.28	5.87	11.39	8.53	3.23	9.76	4.54			-	-
PLACE1000056	6.68	5.77	17.49	7.42	8.83	5.75	5.82	4.2	3.99			-	-
PLACE1000075	12.58	13.3	14.47	10.46	23.08	11.81	12.58	8.38	7.78			-	-
PLACE1000078	16.43	18.48	16.12	10.88	9.95	8.28	13.59	13.6	12.7	**	*	-	-
PLACE1000081	10.24	9.79	10.48	3.99	5.51	5	3.34	5.78	6.89	**	*	-	-
PLACE1000086	24.93	18.79	23.44	11.64	12.45	12.8	14.85	17.83	18.54	**	*	-	-
PLACE1000084	9.77	8.35	20.14	5.83	7.13	8.45	9.11	6.88	4.47			-	-
PLACE1000101	13.24	13.89	14.14	7.47	11.88	8.08	3.54	8.15	5.57	*	**	-	-
PLACE1000121	13.12	18.48	20.42	15.19	28.27	14.52	7.88	10.58	8.77	*		-	-
PLACE1000133	9.91	9.72	12.83	7.9	8.22	8.5	5.09	6.11	8.25	*		-	-
PLACE1000142	9.88	8.89	10.42	4.18	7.46	3.72	7.05	4.74	5.82	*	**	-	-
PLACE1000145	26.84	30.96	24.74	12.94	22.4	10.59	30.36	28.2	28.92	*		-	-
PLACE1000163	7.41	5.19	10.38	3.1	5	4.08	3.16	3.43	3.28	*		-	-
PLACE1000172	27.22	20.45	29.16	10.4	14.2	9.22	8.64	12.45	11.02	**	**	-	-
PLACE1000181	113.9	71.51	73.22	49.28	64.78	42.83	38.78	53.7	55.02			-	-
PLACE1000184	17.72	15.27	19.8	11.79	14.01	13.85	9.11	9.87	10.78	*	**	-	-
PLACE1000185	18.18	14.48	17.72	7.66	7.1	6.34	4.1	6.59	4.27	**	**	-	-
PLACE1000198	9.65	11.81	15.71	11.09	11.55	8.4	6.38	19.65	7.03			-	-
PLACE1000213	8.28	8.05	11.45	4.09	6.38	3.94	3.17	3.29	3.78			-	-
PLACE1000214	9.15	11.17	11.68	8.73	8.08	4.64	3.79	6.02	5.34			-	-
PLACE1000220	25.57	22.54	23.7	11.03	13.14	7.79	10.18	10.81	12.09	**	**	-	-
PLACE1000231	14.42	18.43	23.25	5.64	9.79	6.68	7.39	7.88	9.82	*	*	-	-
PLACE1000236	7.58	6.02	8.68	1.98	8.24	5.81	3.58	10.92	5.78			-	-
PLACE1000245	26.62	25.81	22.88	9.19	16.74	15.35	8.75	9.5	10.2	*	**	-	-
PLACE1000246	45.69	38.58	50.55	14.83	17.18	15.14	8.9	11.47	10.82	**	**	-	-
PLACE1000258	9.85	11.52	11.33	3.18	7.83	3.08	2.45	5.51	4.37	*	**	-	-
PLACE1000288	14.3	14.29	20.78	4.5	8.13	8.1	4.73	7.76	6.77	*	*	-	-
PLACE1000292	22.85	21.9	21.53	9.54	18.83	11.23	12.83	18.24	12.38	*	**	-	-
PLACE1000302	24.34	18.15	25.79	8.82	16.47	8.9	10.89	16.25	19.37	*		-	-
PLACE1000304	9.25	5.15	7.74	5.54	8.18	3.68	3.38	11.18	2.7			-	-
PLACE1000308	15.07	8.68	14.03	8.88	10.28	7.58	8.62	8.08	9.22			-	-
PLACE1000309	50.33	32.12	45.32	24.2	31.82	23.11	35.7	38.12	29.88			-	-
PLACE1000312	7.8	7.45	13.89	4.65	8.18	3.95	7.1	6.14	4.22			-	-
PLACE1000330	37.2	43.13	26.2	23.56	30.84	17.98	20.8	23.35	14.01			-	-
PLACE1000332	19.72	22.93	22.77	9.1	18.25	13.71	15.08	10.86	13.37	*	**	-	-
PLACE1000347	18.65	20.7	25.21	9.08	28.98	17.34	18.08	14.13	12.48	*		-	-
PLACE1000351	4.55	3.12	7.2	2.8	3.17	2.14	6.88	4.21	4.87			-	-

114 / 157

2667

2668

PLACE1000174	4.38	4.25	8.34	4.3	3.67	2.84	1.94	3.1	1.77	*	-
PLACE1000180	16.1	6.81	17.82	9.8	10.53	4.49	5.18	5.02	6.92	*	-
PLACE1000183	6.36	4	6.18	2.37	2.92	2.52	2.89	0.95	0.87	*	-
PLACE1000197	11.59	7.26	11.9	2.79	5.06	5.01	6.09	6.59	6.48	*	-
PLACE1000401	9.04	6.59	7.42	3.23	3.26	3.54	3.28	2.6	2.94	**	-
PLACE1000406	12.08	8.33	14.85	2.59	5.95	3.65	6.86	10.26	6.27	*	-
PLACE1000412	8.51	7.49	7.35	3.42	4.96	3.44	3.84	4.02	2.52	**	-
PLACE1000420	18.93	23.33	27.28	7.24	14.78	9.32	4.07	6.78	6.75	*	-
PLACE1000421	17.83	12.09	14.21	5.67	5.38	4.01	4.24	5.77	6.73	**	-
PLACE1000423	9.91	10.66	9.25	5.17	6.48	3.18	3.29	8.33	2.82	**	-
PLACE1000424	6.41	4.48	6.88	3.52	3.38	1.84	2.89	2.91	2.34	*	-
PLACE1000430	8.9	7.48	10.47	5.49	5	3.45	3.57	3.09	3.55	*	-
PLACE1000433	6.1	6.1	7.11	1.95	4.44	2.14	2.34	1.88	2.11	*	-
PLACE1000435	28.95	26.63	24.32	14.09	14.79	9.09	16.84	11.65	12.23	**	-
PLACE1000437	13.7	9.76	13.69	6.32	11.96	9.08	8.89	6.76	4.42	*	-
PLACE1000442	4.21	4.33	6.54	1.97	3.93	1.28	2.62	4.3	2.3		-
PLACE1000444	6.39	8.02	6.47	3.23	8.73	7.32	3.84	5.13	5.23		-
PLACE1000453	8.24	8.08	7.4	2.64	6.34	3.86	5.87	6.05	5.1	*	-
PLACE1000456	9.14	8.93	7.88	4.6	6.83	3.55	4.93	5.65	3.15	*	-
PLACE1000485	29.59	27.02	24.71	10.68	16.28	9.53	7.25	7.03	4.81	**	-
PLACE1000481	10.19	8.45	9.18	6.24	5.2	4.46	5.55	2.41	2.95	**	-
PLACE1000492	14.54	13.99	15.18	6.98	17.38	5.81	6.24	4	6.05	**	-
PLACE1000508	8.74	6.29	6.02	2.48	6.75	1.48	4.68	3.01	3.04	*	-
PLACE1000512	4.85	3.74	3.65	2.37	3	3.01	2.27	2.44	1.75	*	-
PLACE1000540	12.96	8.47	7.79	4.5	9.85	7.03	5.75	9.5	3.54		-
PLACE1000541	10.18	16.98	11.44	6.25	17.26	10.37	7.65	8.38	9.85		-
PLACE1000545	13.1	11.96	11.32	7.18	10.25	10.48	6.08	5.2	6.26	**	-
PLACE1000547	8.49	6.52	7.68	5.49	18.6	14.98	5.43	2.62	3.64	*	-
PLACE1000560	22.37	15.31	18.14	8.38	18.85	7.68	6.26	3.35	3.9	*	-
PLACE1000562	6.22	7.18	8	4.82	4.33	3.02	4.18	2.13	2.7	**	-
PLACE1000564	17.41	14.24	11.25	10.42	18.44	10.34	8.72	6.17	6.09	*	-
PLACE1000583	11.44	8.04	7.49	4.72	4.9	3.91	5.24	5.95	5.28	*	-
PLACE1000587	7.15	6.59	3.7	3.14	3.88	4.15	3.78	3.64	4.07		-
PLACE1000588	18.91	11.92	9.8	5.65	7.81	8.38	8.6	5.83	6.87		-
PLACE1000596	11.7	13.68	11.65	8.47	13.84	9.08	10.58	6.53	9.44		-
PLACE1000599	9.78	11.08	9.84	6.95	18.5	6.09	4.81	2.43	3.18	**	-
PLACE1000605	16.65	12.34	15.55	9.49	9.68	6.14	6.32	3.07	5.49	*	-
PLACE1000610	13.13	11.67	11.01	8.74	7.05	4.41	9.89	6.2	4.65	*	-
PLACE1000611	44.24	42.43	32.52	28.13	21.67	19.73	63.55	29.23	50.7		-
PLACE1000626	7.18	6.35	6.83	3.54	3.14	2.93	5.25	5.57	5.32	**	-
PLACE1000631	7.76	6.07	5.81	4.09	6.95	4.33	4.51	3.72	5.54	*	-
PLACE1000635	9.18	4.57	3.73	3.4	5.99	3.07	2.89	4.22	5.33		-
PLACE1000653	13.52	15.38	9.28	7.56	11.63	7.5	6.22	4.09	7.58	*	-
PLACE1000656	16.45	24.27	20.78	6.6	13.98	8.95	13.7	10.63	9.13	*	-
PLACE1000683	17.59	14.23	11.72	6.71	8	7.43	15.75	11.07	18.5	*	-
PLACE1000706	22.58	27.08	16.29	14.32	16.15	9.77	10.6	7.13	11.87	*	-
PLACE1000712	15.91	18.32	12.16	10.41	18.03	9.89	15.21	13.38	14.12		-
PLACE1000716	9.51	6.38	10.32	4.38	8.28	5.91	5.52	6.18	6.28		-
PLACE1000740	62.65	63.87	75.82	66.13	68.56	67.41	54.02	47.28	39.42	*	-
PLACE1000748	21.69	20.13	21.77	15.81	19.87	14.26	18.35	17.23	18.62	*	-
PLACE1000749	6.88	6.77	9.49	5.21	7.14	4.97	2.01	3.41	3.49	**	-
PLACE1000751	5.85	6.89	7.78	3.42	4.79	3.39	2.03	3.31	2.82	*	-
PLACE1000755	17.27	23.58	23.84	13.28	21.84	9.25	15.55	10.98	9.19	*	-
PLACE1000769	3.3	5.09	3.02	3.65	5.58	3.4	3.76	2.68	2.93		-
PLACE1000778	43.25	43.09	44.83	22.77	18.03	27.54	20.87	24.08	21.65	**	-
PLACE1000785	31.35	17.11	31.03	9.35	9.07	8.25	11.86	6.82	7.46	*	-
PLACE1000786	5.99	3.16	7.12	2.4	4.01	1.88	2.45	4.94	6.92	*	-
PLACE1000793	62.95	70.84	66.12	74.58	80.1	68.58	59.03	72.61	48.69		-
PLACE1000795	170.82	165.32	165.81	123.39	175.42	137.1	104.41	82.02	95.27	**	-
PLACE1000798	34.34	30.96	28.31	19.04	23.22	14.81	22.05	22.97	20.25	*	-
PLACE1000812	18.41	17.23	19.15	11.88	12.41	10.24	15.41	13.57	12.8	**	-
PLACE1000823	25.51	22.51	19.45	15.2	17.94	10.61	13.45	15.88	12.98	*	-
PLACE1000825	8.78	4.9	7.94	5.92	3.89	3.88	4.94	3.55	4.1	*	-
PLACE1000838	37.15	32.03	40.12	17.88	18.59	17.35	19.92	15.21	15.97	**	-

115 / 157

2669

2670

PLACE1000841	7.87	7.94	8.8	7.6	12.23	10.86	5.33	4.89	4.56	**	-	-
PLACE1000842	18.04	20.1	24.44	8.07	13.38	10.89	7.02	9.18	7.59	*	-	-
PLACE1000849	5.43	7.23	9.29	4.99	5.8	4.19	3.26	3.86	2.16	*	-	-
PLACE1000856	209.88	229.07	253.03	209.39	258.66	183.75	147.93	119.29	119.6	**	-	-
PLACE1000863	16.12	16.52	27.27	10.41	12.7	10.18	13.62	16.56	12.54	*	-	-
PLACE1000876	16.98	21.81	24.48	11.69	18.7	12.24	19.45	16.78	20.21	*	-	-
PLACE1000889	225.37	148.25	253.81	108.96	112.25	68.21	130.08	130.64	120.26	*	-	-
PLACE1000907	11.9	9.88	11.62	4.52	6.53	6.58	4.58	7.2	4.58	**	**	-
PLACE1000909	219.81	187.78	224.87	128.92	144.43	97.69	289.39	207.09	231	**	-	-
PLACE1000912	4.53	4.31	6.11	1.95	4.16	2.65	2.78	1.25	1.4	*	-	-
PLACE1000914	13.65	11.52	18.16	6.78	8.53	7.59	5.22	6.53	2.91	*	-	-
PLACE1000918	13.18	11.98	16.82	5.88	8.64	8.52	8.09	14.67	8.16	*	-	-
PLACE1000927	6.93	7.22	10.64	4.38	6.32	3.11	4.55	3.47	4.34	*	-	-
PLACE1000931	8.5	7.15	10.84	7.43	6.57	4.4	3.12	4.27	3.39	**	-	-
PLACE1000944	12.44	7.29	10.38	5.9	6.11	4.83	6.15	8.79	6.0	*	-	-
PLACE1000948	15.03	15	16.78	7.58	11.29	7.62	5.6	8.31	6.83	**	**	-
PLACE1000958	5.27	4.99	7.38	2.51	5.7	3.89	1.85	6.23	4.43	*	-	-
PLACE1000972	8.38	5.92	10.49	3.85	7.53	4.25	3.77	5.68	4.19	*	-	-
PLACE1000977	11.28	8.12	15.95	5.04	5.25	7.02	2.5	10.48	3.14	*	-	-
PLACE1000979	13.89	7.88	9.88	3.34	7.43	5.36	4.42	5.83	8.8	*	-	-
PLACE1000986	30.12	30.53	34.37	25.05	28	23.07	15.94	15.63	18.52	*	**	-
PLACE1000987	19.26	23.97	21.74	13.41	21.5	10.81	16.26	13.02	12.47	*	-	-
PLACE1001000	4.37	7.15	10.87	3.42	3.04	2.14	2.69	2.93	3.43	*	-	-
PLACE1001007	6.61	2.96	5.44	2.81	6.06	3.04	2.78	3.89	2.01	*	-	-
PLACE1001010	9.48	6.18	10.43	5.02	5.52	3.2	5.18	3.73	3.91	*	-	-
PLACE1001015	7.99	5.3	8.09	6.19	7.65	4.6	3.28	3.31	2.3	*	-	-
PLACE1001016	17.51	11.26	16.44	6.81	7.04	6.42	3.95	3.27	2.38	*	**	-
PLACE1001022	14.42	12.9	14.76	4.87	7.37	5.58	7.11	8.21	7.4	**	**	-
PLACE1001024	11.18	12.76	12	5.14	19.39	4.2	4.43	3.88	2.99	**	-	-
PLACE1001036	105.59	109.4	58.51	52.8	114.22	89.89	113.54	86.02	137.21	*	-	-
PLACE1001038	9.63	8.88	10.53	4.88	5.51	5.33	2.98	5.92	7.98	**	-	-
PLACE1001048	7.22	3.94	6.41	5.19	5.21	3.49	3.25	3.01	2.83	*	-	-
PLACE1001054	17.12	18.15	23	21.95	14.11	11.96	6.57	6.82	6.81	**	-	-
PLACE1001062	7.89	5.88	7.49	3.03	6.18	4.88	5.8	1.77	1.91	*	-	-
PLACE1001063	11.62	13.23	13.69	8.34	8.29	7.45	3.15	3.88	2.78	**	**	-
PLACE1001076	17.48	15.27	22.58	9.61	12.22	10.09	9.5	6.99	10.2	*	-	-
PLACE1001081	8.04	6.12	10.01	2.34	3.8	3.43	2.11	2.93	3.84	*	-	-
PLACE1001088	1077.5	1434.9	897.35	789.08	838.84	874	835.48	1189	1173.1	*	-	-
PLACE1001082	41.3	35.7	38.59	14.58	20.22	12.39	28.15	30.94	33.71	**	-	-
PLACE1001098	27.38	22.29	18.87	5.94	11.11	13.11	8.05	8.38	7.88	*	**	-
PLACE1001100	12.04	12.72	14.57	7.69	8.68	8.84	7.95	8.39	9.96	**	-	-
PLACE1001104	10.26	8.03	11.29	7.1	4.81	4.72	5.84	4.16	5.12	*	-	-
PLACE1001114	19.87	21.11	13.11	5.44	7.05	8.22	2.52	3.44	2.49	*	**	-
PLACE1001118	18.04	16.17	18.54	9.49	13.49	10.14	5.69	6.26	3.99	**	-	-
PLACE1001123	11.13	9.28	10.37	3.01	8.71	7.98	2.59	2.22	5.96	**	-	-
PLACE1001128	9.63	8.38	10.4	4.16	7.83	10.1	11.4	6.43	4.95	*	-	-
PLACE1001144	22.24	18.89	18.24	5.11	10.59	8.11	16.88	14.03	17.77	**	-	-
PLACE1001147	9.73	14.79	11.54	7.5	5.73	6.98	3.33	4.01	4.07	*	**	-
PLACE1001148	16.26	12.48	13.41	8.25	12.02	6.54	11.91	10.71	11.91	*	-	-
PLACE1001159	5.83	2.18	9.14	5.55	4.85	2.75	3.37	5.16	1.28	*	-	-
PLACE1001168	6.64	5.38	5	2.78	4.75	2.95	4.38	2.12	2.38	*	-	-
PLACE1001171	17.87	14.63	16.08	9.77	8.8	9.82	9.4	7.25	9.58	**	**	-
PLACE1001183	24.74	23.48	20.67	15.23	15.9	11.49	12.71	7.91	11.38	**	**	-
PLACE1001186	8.91	7.02	8.02	4.21	7.14	4.11	5.58	3.21	2.94	*	**	-
PLACE1001201	16.85	15.12	18.88	10.04	10.01	9	9.33	10.2	8.94	**	**	-
PLACE1001229	15.21	12.81	13.94	8.57	11.28	12.88	8.81	8.47	8.63	**	-	-
PLACE1001231	13.31	19.61	12.2	8.58	26.11	14.91	13.39	13.43	8.59	*	-	-
PLACE1001238	8.18	6.8	5.78	3.46	5.45	5.07	3.57	1.84	3.83	**	-	-
PLACE1001241	7.81	12.55	8.77	4.58	8.8	2.29	6.75	1.69	5.45	*	-	-
PLACE1001242	15.16	9.43	9.57	6.93	8.62	5.05	5.42	2.88	4.67	*	-	-
PLACE1001247	18.19	25.95	17.43	7.95	13.13	8.09	9.19	7.63	8.42	*	-	-
PLACE1001250	27.19	18.6	15.62	14.1	21.99	16.23	13.87	8.35	7.96	*	-	-
PLACE1001257	5.07	4.07	3.3	8.21	3.6	2.05	3.84	3.01	2.82	*	-	-
PLACE1001272	11.61	5.34	6.33	4.73	5.73	6.35	5.18	6.02	6.39	*	-	-

2671

2672

PLACE1001270	20.66	18.57	13.78	8.04	11.1	8.58	6.47	4.72	6	*	**	-	-
PLACE1001280	13.73	13.22	15.63	11.44	10.47	6.22	11.2	7.22	13.37	*	**	-	-
PLACE1001294	6.1	5.02	7.03	2.50	6.24	4.37	4.18	2.25	4.38	*	**	-	-
PLACE1001295	9.52	6.34	5.73	4.11	5.38	4.8	4.68	3.72	5.18	*	**	-	-
PLACE1001300	110.16	138.11	98.73	55.88	77.8	89.78	127.88	93.28	123.22	*	**	-	-
PLACE1001304	22.83	23.38	19.82	13.09	11.18	12.58	8.64	8.3	14.22	*	**	-	-
PLACE1001311	22.83	29.08	18.33	13.63	11.98	8.81	18.15	14.72	19.17	*	**	-	-
PLACE1001323	6.15	4.73	4.73	2.24	3.77	2.07	2.58	2.19	2.91	*	**	-	-
PLACE1001325	6.51	4.28	2.68	3.73	4.01	0.79	3.09	1.19	1.69	*	**	-	-
PLACE1001340	12.71	9.9	7.81	5.83	10.17	3.62	8.49	5.49	7.3	*	**	-	-
PLACE1001344	7.28	5.81	8.3	3.22	4.87	2.35	5.25	3.25	5.47	*	**	-	-
PLACE1001351	10.82	9.34	9.86	4.54	4.04	2.83	5.88	5.94	6.17	*	**	-	-
PLACE1001356	18.48	18.32	12.21	7.88	10.81	8	10.02	6.25	6.08	*	**	-	-
PLACE1001377	10.72	11.21	8.12	8.14	11.01	6.18	7.03	5.07	5.96	*	**	-	-
PLACE1001383	8.47	7.74	7.34	7.01	7.31	9.18	4.38	4.51	5.57	*	**	-	-
PLACE1001384	5.51	3.13	4.87	2.28	4.64	2.12	1.25	1.77	2.28	*	**	-	-
PLACE1001387	3.71	2.66	5.28	2.18	4.13	3.15	1.31	2.35	1.75	*	**	-	-
PLACE1001395	2.97	1.83	2.85	0.8	3.39	0.84	2.28	2.27	0.83	*	**	-	-
PLACE1001399	43.48	38.85	39.47	25.54	38.02	28.91	11.4	22.35	12.37	*	**	-	-
PLACE1001401	8.85	12.13	13.82	4.1	7.84	5.28	5.15	6.58	4.43	*	**	-	-
PLACE1001407	8.42	8.85	8.88	3.19	5.6	3.04	3.23	2.38	2.93	*	**	-	-
PLACE1001412	4.65	3.97	3.58	2.52	8.48	2.63	3.1	2.07	4.54	*	**	-	-
PLACE1001414	10.05	7.04	8.23	6.28	7.99	5.53	4.37	7.32	4.41	*	**	-	-
PLACE1001416	18.05	14.76	21.75	8.21	15.69	7.54	12.88	14.94	10.13	*	**	-	-
PLACE1001433	10.35	9.63	10.38	6.47	8.28	6.35	4.33	6.8	5.78	*	**	-	-
PLACE1001440	15.78	8.2	12.08	9.87	10.51	7.97	4.11	5.02	3.17	*	**	-	-
PLACE1001456	5.95	5.2	7.6	4.32	4.51	3.88	3.45	6.83	4.09	*	**	-	-
PLACE1001484	7.17	7.13	12.08	4.48	13.28	4.47	2.82	3.59	4.74	*	**	-	-
PLACE1001488	5.87	5.89	7.51	4.49	7.19	3.15	4.98	3.88	5.45	*	**	-	-
PLACE1001484	1.93	2.72	3.26	3.17	5.24	2.3	3.27	1.7	4.61	*	**	-	-
PLACE1001500	5.55	5.43	7.47	3.4	4.88	4.91	3.43	5.41	4.22	*	**	-	-
PLACE1001502	10.13	8.73	10.38	4.95	5.09	3.9	4.24	7.4	4.55	*	**	-	-
PLACE1001503	18.94	15.22	23.11	8.82	14.8	9.42	9.19	12.06	11.89	*	**	-	-
PLACE1001505	14.33	11.5	15.08	6.94	9.91	8.35	5.05	5.21	4.29	*	**	-	-
PLACE1001513	9.76	7.84	14.27	4.51	12.01	6.72	4.02	8.18	4.23	*	**	-	-
PLACE1001516	12.33	10.13	14.29	8.5	8.85	7.91	5.4	6.92	6.78	*	**	-	-
PLACE1001517	22.78	28.83	26.65	15.8	22.43	12.48	25.55	23.34	19.3	*	**	-	-
PLACE1001523	4.86	4.2	4.25	1.48	3.84	2.44	3.81	4.37	5.02	*	**	-	-
PLACE1001526	15.63	8.8	12.83	5.9	5.99	8.7	6.38	8.18	11.68	*	**	-	-
PLACE1001534	13.26	10.83	12.93	4.83	7.68	4.05	5.45	22.34	4.41	*	**	-	-
PLACE1001536	12.35	10.07	11.17	5.95	10.51	7.42	7.47	10.57	7.69	*	**	-	-
PLACE1001545	7.78	9.31	13.04	5.48	7.28	3.94	2.6	3.19	3.02	*	**	-	-
PLACE1001551	11.64	13.9	20.49	7.15	8.98	9.21	7.74	8.57	7.78	*	**	-	-
PLACE1001564	6.33	7.01	11.33	5.54	6.02	8.32	3.9	3.45	8.93	*	**	-	-
PLACE1001570	8.81	7.17	13.9	5.81	7.77	3.89	5.34	9.04	4.82	*	**	-	-
PLACE1001571	30.18	28.44	29.75	18.26	25.49	9.78	12.94	16.78	20.12	*	**	-	-
PLACE1001595	48.16	53.02	65.48	28.84	28.86	28.28	20.83	17.19	22.48	*	**	-	-
PLACE1001602	17.83	20.88	26.68	9.8	15.68	12.03	25.23	43.22	27.97	*	**	-	-
PLACE1001603	18	11.85	13.84	6.25	7.5	8.17	10.37	25.06	14.43	*	**	-	-
PLACE1001608	313.79	262.25	289.49	248.62	184.68	212.51	255.55	258.42	224.33	*	**	-	-
PLACE1001610	34.83	33.4	50.14	21.2	18.91	21.09	18.25	16.43	15.58	*	**	-	-
PLACE1001611	18.58	14.57	13.52	8.27	15.57	8.52	5.98	3.97	4.4	*	**	-	-
PLACE1001629	15.88	14.27	16.52	8.82	12.73	6.97	9.05	9.4	8.15	*	**	-	-
PLACE1001632	15.8	14.68	11.95	5.15	11.43	6.75	4.77	3.59	2.95	*	**	-	-
PLACE1001634	10.25	9.24	12.21	4.71	7.41	4.15	3.93	4.97	4.83	*	**	-	-
PLACE1001637	29.75	28.67	38.48	18.79	18.46	16.75	23.41	26.78	22.84	*	**	-	-
PLACE1001640	28.95	18.03	24.89	15.83	21.97	12.71	8.91	11.82	9.32	*	**	-	-
PLACE1001655	8.24	5.58	6.83	4.09	4.25	2.92	2.13	3.7	1.34	*	**	-	-
PLACE1001672	8.94	18.15	13.65	4.81	11.29	4.01	5.1	4.86	14.47	*	**	-	-
PLACE1001676	38.58	47.75	47.96	23.16	26.28	21.98	31.5	32.99	28.58	*	**	-	-
PLACE1001683	7.54	6.2	11.82	2.18	4.9	3.68	2.93	3.43	3.72	*	**	-	-
PLACE1001691	10.4	6.5	11.03	4.05	13.34	5.07	4.28	5.31	2.44	*	**	-	-
PLACE1001692	3.74	6.24	4.95	3.58	3.97	3.4	3.88	3.22	2.42	*	**	-	-
PLACE1001705	11.04	7.84	13.58	10.64	5.83	6.51	5.2	5.34	3.62	*	**	-	-

117 / 157

2673

2674

PLACE1001718	13.89	8.26	12.85	5.88	7.61	3.53	5.17	4.09	5.84	*	**	-	-
PLACE1001720	28.23	23.88	21.88	17.41	18.42	17.41	10.88	9.34	8.72	*	*	-	-
PLACE1001728	21.97	14.68	15.38	7.87	10.67	10.96	10.39	5.48	5.25	*	*	-	-
PLACE1001729	7.28	8.78	9.32	4.24	3.8	3.29	3.98	2.03	2.64	**	**	-	-
PLACE1001739	18.08	18.75	18.08	11.34	16.38	11.72	10.55	8.28	8.75	*	**	-	-
PLACE1001740	60.6	61.68	51.85	52.21	47.26	48.39	41.09	28.81	37.51	**	**	-	-
PLACE1001745	155.43	121.46	146.04	80.89	92.15	78.37	106.17	102.98	100.25	**	*	-	-
PLACE1001748	12.07	9.53	11.89	5.08	6.32	6.38	8.4	7.42	7.33	**	**	-	-
PLACE1001748	20.09	15.85	19.67	8.98	18.04	8.77	6.58	3.28	8.51	*	*	-	-
PLACE1001753	8	5.61	7.17	2.42	2.42	2.08	3.13	1.99	2.1	**	**	-	-
PLACE1001760	16.81	13.89	14.08	5.89	8.99	11.7	8.88	6.91	7.81	*	**	-	-
PLACE1001767	7.75	5.93	9.84	3.14	5.58	2.6	2.92	2.72	1.2	*	**	-	-
PLACE1001771	6.88	8.78	6.6	6.1	22.6	7.92	6.68	3.61	1.79	*	*	-	-
PLACE1001775	10.25	8.72	8.89	5.62	4.98	4.94	5.79	4.51	5.33	**	**	-	-
PLACE1001777	55.87	33.24	48.38	18.69	28.37	22.71	28.45	22.89	29.83	*	*	-	-
PLACE1001781	7.82	4.99	6.26	2.61	3.65	3.27	2.46	2.47	3	*	*	-	-
PLACE1001783	8.67	3.54	4.17	1.75	1.71	2.1	2.79	1.87	2.04	*	*	-	-
PLACE1001788	10.13	7.03	8.19	5.28	7.53	4.48	4.3	2.57	2.1	*	*	-	-
PLACE1001788	9.62	8.2	7.79	2.13	2.02	3.87	3.97	1.93	2.04	**	**	-	-
PLACE1001795	12.86	13.57	11.81	3.78	6.82	4.77	2.63	4.65	2.65	**	**	-	-
PLACE1001799	15.19	12.46	11.73	6.44	10.22	5.84	11.68	8.72	12.14	*	*	-	-
PLACE1001810	19.9	17.99	15.3	8.61	18.81	15.9	19.42	15.41	19.52	**	**	-	-
PLACE1001817	8.2	10.03	10.24	5.73	12.6	6.09	4.49	4.8	5.41	*	*	-	-
PLACE1001821	8.12	9.65	9.24	5.34	6.81	5.22	5.09	4.7	1.49	**	**	-	-
PLACE1001836	14.74	10.74	8.7	7.69	8.81	4.93	6.55	5.02	5	*	*	-	-
PLACE1001844	8.12	5.11	7.16	1.98	5.82	2.65	3.18	2.67	1.5	*	*	-	-
PLACE1001845	12.24	11.82	8.77	7.69	7.48	3.74	4.84	2.2	2.42	**	**	-	-
PLACE1001858	9.08	7.62	7.23	4.18	3.93	3.22	7.86	3.49	3.91	**	**	-	-
PLACE1001869	5.88	6.08	4.84	3.37	6.33	2.79	3.14	2.33	2.54	**	**	-	-
PLACE1001890	14.71	9.13	7.02	7.27	10.27	7.92	6.48	7.41	7.67	*	*	-	-
PLACE1001897	21.83	13.66	12.13	7	10.58	7.47	6.17	7.42	7.81	*	*	-	-
PLACE1001902	25.9	24.38	23.51	11.89	18.02	13.01	23.32	19.87	28.51	*	*	-	-
PLACE1001904	4.8	4.58	4.71	4.2	4.02	2.48	4	1.2	3.18	*	*	-	-
PLACE1001907	11.99	11.09	12.77	4.36	7.42	4.55	5.4	2.81	8.4	**	**	-	-
PLACE1001910	17.55	13	11.78	10.79	10.88	3.78	8.75	3.34	8.51	*	*	-	-
PLACE1001912	16.89	20.78	18.01	10.56	10.89	7.91	12.99	8.39	12.1	**	*	-	-
PLACE1001918	10.35	11.16	9.4	5.64	8.81	4.78	8.14	4.97	7.28	*	*	-	-
PLACE1001920	3.62	1.84	1.09	2.08	2.78	2.12	0.89	2.37	2.67	*	*	-	-
PLACE1001928	33.91	29.31	26.67	16.07	24.12	10.81	21.1	17.01	25.48	*	*	-	-
PLACE1001930	13.35	8.41	9.7	5.18	8.65	8.14	5.7	4.65	8.73	*	*	-	-
PLACE1001949	83.24	64.21	68.63	50.5	75.89	68.37	92.93	85.41	92.12	*	*	-	-
PLACE1001953	8.04	5.93	10.84	4.58	6.1	3.95	4.33	6.08	5.07	*	*	-	-
PLACE1001968	8.52	5.51	4.9	2.62	4.38	2.04	3.84	2.43	3.2	*	*	-	-
PLACE1001974	16.8	19.43	17.28	9.77	10.54	5.8	6.98	5.48	6.49	**	**	-	-
PLACE1001981	11.42	11.82	8.18	7.19	8.78	5.34	7.9	6.15	5.59	*	*	-	-
PLACE1001983	9.22	8.71	12.62	5.64	7.95	6.32	7.39	5.69	5.5	*	*	-	-
PLACE1001989	27.47	22.77	28.46	13.3	18.78	12.49	12.35	15.17	10.35	*	**	-	-
PLACE1002084	8.73	7.48	8.33	6.79	8.77	5.49	8.57	8.13	4.33	*	*	-	-
PLACE1002088	10.54	8.97	12.2	6.77	11.48	10.25	3.74	7.33	3.11	*	*	-	-
PLACE1002016	12.37	10.1	13.16	6.64	10.68	6.15	7.89	14.89	9.22	*	*	-	-
PLACE1002044	18.51	15.04	21.25	6.21	11.74	9.02	8.6	10.2	12.21	*	*	-	-
PLACE1002046	3.16	4.19	5.33	2.48	8.58	3.13	3.41	1.95	2.73	*	*	-	-
PLACE1002052	4.29	2.34	2.47	1.89	4.94	1.55	4.09	3.83	3.62	*	*	-	-
PLACE1002066	4.6	3.49	6.01	2.29	3.79	2.93	3.54	3.65	2.92	*	*	-	-
PLACE1002072	6.4	5.64	8.81	3.88	5.18	4.7	3.85	6.12	2.07	*	*	-	-
PLACE1002073	12.89	11.8	16.18	10.66	11.47	7.97	5.88	5.64	4.74	**	**	-	-
PLACE1002080	10.63	22.97	23.1	12.21	18.28	13.9	13.6	15	15.2	*	*	-	-
PLACE1002081	23.13	27.18	27.3	14.91	18.08	14.88	16.14	20.45	20.82	**	*	-	-
PLACE1002090	11.49	14.13	12.79	4.54	7.24	4.65	4.29	3.51	5.75	**	**	-	-
PLACE1002095	5.45	5.54	4.45	5.08	11.47	7.41	4.99	5.27	7.8	*	*	-	-
PLACE1002102	8.22	13.54	14.09	8.11	14.1	7.74	9.04	18.2	14.08	*	*	-	-
PLACE1002109	17.45	10.93	19.35	12.82	7.15	7.81	9.22	12.81	8.47	*	*	-	-
PLACE1002115	6.48	5.79	9.09	2.86	4.39	3.54	3.32	3.75	3.44	*	*	-	-
PLACE1002119	3.98	5.58	7.03	2.87	6.47	3.2	3.72	4.12	4.62	*	*	-	-

118 / 157

2675

2676

PLACE1002140	8.54	9.5	15.02	4.83	8.88	8.88	2.91	8.38	4.28	*	*	-
PLACE1002150	27.18	33.81	30.05	18.28	25.09	23.98	12.7	14.95	15.29	*	**	-
PLACE1002153	173.25	217.95	284.37	422.44	424.65	291.99	262.39	242.27	220.74	*	+	-
PLACE1002157	7.42	5.99	8.84	3.18	6.59	2.62	3.59	2.5	5.34	*	*	-
PLACE1002163	9.5	10.33	17.61	4.81	9.35	4.71	6.81	8.07	8.48	*	*	-
PLACE1002168	408.18	407.82	542.62	187.65	60.86	161.78	377.56	231.11	429.49	**	-	-
PLACE1002170	23.07	18.88	22.82	9.58	14.95	10.12	4.39	8.88	5.14	*	**	-
PLACE1002171	118.3	86.27	82.6	107.58	135.49	98.73	60.2	58.88	56.83	*	*	-
PLACE1002180	12.53	12.02	17.4	11.66	15.77	12.8	11.81	13.11	10.84	*	*	-
PLACE1002184	76.94	90.34	67.89	42.12	70.88	48.23	20.87	27.8	29.07	**	-	-
PLACE1002200	5.84	6.44	11.68	2.73	4.37	3.29	2.8	2.86	3.48	*	*	-
PLACE1002205	5.47	5.6	9.99	2.28	6.05	2.28	3.51	2.6	1.9	*	-	-
PLACE1002213	11	9.88	7.88	3.37	8.5	3.33	5.73	2.26	4.33	*	-	-
PLACE1002219	30.67	24.88	33.1	11.7	18.03	17.11	5.85	10.58	12.01	*	**	-
PLACE1002227	5.38	6.1	6.51	3.19	5.95	3.48	3.48	8.52	2.37	*	-	-
PLACE1002253	7.93	4.48	10.74	3.51	5.52	2.84	4.18	3.78	3.24	*	-	-
PLACE1002256	10.19	4.49	11.42	3.86	5.44	3.17	1.81	4.48	5.21	*	-	-
PLACE1002259	6.68	9.14	16.04	2.32	8.56	3.26	2.76	5.45	13.81	*	-	-
PLACE1002285	15.21	20.2	16.69	10.84	18.4	10.19	8.14	7.95	6.12	**	-	-
PLACE1002301	11.84	7.92	13.87	3.5	6.75	4.81	3.98	8.04	4.04	*	-	-
PLACE1002310	9.3	8.21	7.36	1.68	5.25	3.58	3.57	2.81	3.34	*	**	-
PLACE1002311	6.47	4.99	7.42	3.5	6.65	2.75	2.49	2.97	1.32	*	-	-
PLACE1002319	23.43	17.43	19.83	7.68	14.77	10.74	10.78	13.23	11.96	*	-	-
PLACE1002329	7	4.89	6.25	3.08	3.73	2.65	1.68	2.85	1.64	*	**	-
PLACE1002333	16.35	14.3	13.78	9.29	11.1	10.63	8.49	6.32	5.51	**	-	-
PLACE1002342	14.18	15.16	20.35	5.96	11.78	6.24	4.58	8.15	6.84	*	**	-
PLACE1002343	9.54	8.66	16.84	3.45	4.23	3.21	2.88	1.85	3.18	*	-	-
PLACE1002355	8.47	11.34	12.23	4.32	8.27	5.65	2.98	2.78	4.19	*	-	-
PLACE1002358	6.25	5.01	7.85	2.33	4.09	1.68	2.63	1.27	2.04	*	**	-
PLACE1002359	94.59	64.3	78.28	42.47	28.68	29.68	51.17	53.76	59.14	**	-	-
PLACE1002374	11.84	7.08	11.73	4.84	7	4.57	3.23	3.62	3.66	*	-	-
PLACE1002376	10.64	6.42	7.57	5.56	5.2	4.83	8.79	6.04	8.82	*	-	-
PLACE1002379	14.26	10.82	12.17	8.95	8.14	6.98	5.34	4.44	3.8	**	**	-
PLACE1002386	7.89	3.52	7.76	1.91	2.26	2.95	2.4	2.03	2.94	*	-	-
PLACE1002395	17.68	20.65	22.77	6.45	8.6	7.04	4.3	2.63	2.39	**	**	-
PLACE1002399	9.59	8.59	10	4	5.08	4.5	2.83	2.8	3.48	**	**	-
PLACE1002407	14.84	17.54	12.15	4.78	9.29	6.64	5.32	3.95	5.87	*	**	-
PLACE1002433	18.89	14.68	17.11	8	12.47	8.81	12.74	12.61	18.41	*	-	-
PLACE1002437	6.94	5.04	7.88	4.77	3.72	3.8	1.87	3.5	4.81	*	-	-
PLACE1002438	8.47	9.78	8.93	8.11	8.03	2.52	2.69	3.74	4.55	*	**	-
PLACE1002446	61.22	60.91	53.35	35.08	40.23	34.18	52.04	54.81	48.63	**	-	-
PLACE1002447	58.08	38.25	48.75	30.92	38.93	28.38	28.38	23.5	18.47	**	-	-
PLACE1002458	8.68	4.18	8.14	2.58	3.07	3.08	3.45	3.27	1.68	*	-	-
PLACE1002462	20.62	21.36	20.69	8.83	13.93	7.63	14.57	11.49	17.87	**	*	-
PLACE1002485	6.3	10.21	4.45	2.37	6.62	4.25	3.41	1.81	3.99	*	-	-
PLACE1002477	8.9	10.86	8.01	5.33	8.28	3.73	2.36	3.29	4.22	**	-	-
PLACE1002493	10.23	12.91	13.31	8.45	8.55	11.89	12.5	8.8	14.98	*	-	-
PLACE1002497	5.44	5.3	4.58	4.99	2.83	4.56	5.81	1.5	1.66	*	-	-
PLACE1002499	9.23	8.42	6.03	5.64	3.94	3.18	4.77	3.01	2.42	*	-	-
PLACE1002500	28.78	23.91	22.93	13.69	10.19	9.79	16.45	7.7	12.71	**	-	-
PLACE1002514	9.83	6.14	8.08	4.34	6.18	2.49	5.4	1.41	3.42	*	-	-
PLACE1002518	67.39	61.98	51.39	31.15	38.98	29.56	53.37	39.47	40.85	**	-	-
PLACE1002529	7.97	7.12	4.14	3.92	5.36	3.52	3.58	1.82	2.65	*	-	-
PLACE1002532	14.35	10.84	8.84	6.09	12.13	8.95	3.37	3.8	3.7	**	-	-
PLACE1002535	6.92	8.97	5.35	4.94	5.47	4.78	4.62	5.38	3.19	*	-	-
PLACE1002536	9.3	12.32	10.38	5.35	4.64	5.41	7.12	5.91	4.69	**	*	-
PLACE1002537	7.11	8.85	6.37	5.89	5.45	6.3	4.25	3	3.53	**	-	-
PLACE1002539	7.92	6.45	7.28	2.97	5.46	7.97	2.07	2.63	4.82	*	-	-
PLACE1002547	14.65	14.48	12.7	9.78	11.89	6.33	5.01	4.47	6.38	**	-	-
PLACE1002571	27.85	40.71	17.49	16.84	14.71	12.01	13.72	7.48	9.31	*	-	-
PLACE1002578	11.7	10.9	10.03	8.42	11.59	6.31	11.22	5.24	9.68	*	-	-
PLACE1002583	12.45	8.98	8.93	4.7	6.58	6.37	8.82	7.31	10.13	*	-	-
PLACE1002591	14.95	9.64	30.55	14.89	14.5	1.99	10.65	10.99	7.37	*	-	-
PLACE1002598	134.77	135.2	115.7	105.05	127.67	81.8	77.51	75.71	89.77	**	-	-

119 / 157

2677

2678

PLACE1002604	8.27	3.87	5.87	4.56	4.62	4.03	4.03	2.97	4.32										
PLACE1002612	7.74	6.07	5.48	4.79	5.04	4.28	2.87	1.84	4.68										
PLACE1002625	8.88	7.38	5.28	3.81	6.23	4.29	4.61	2.42	5.37										
PLACE1002638	10.71	6.82	9.31	4.39	4.89	3.67	4.88	2.91	4.12	*	*	-	-	-	-	-	-	-	-
PLACE1002655	20.05	17.22	10.12	8.15	9.62	8.61	4.45	4.4	4.69	*	*	-	-	-	-	-	-	-	-
PLACE1002665	2.85	2.11	1.16	1.34	2.18	0.77	1.38	1.74	1.47										
PLACE1002685	40.11	36.95	37.09	21.27	32.67	18.25	35	28.12	33.61	*	*	-	-	-	-	-	-	-	-
PLACE1002692	10.4	8.99	7.49	3.22	6.32	3.77	7.81	3.78	6.83	*	*	-	-	-	-	-	-	-	-
PLACE1002714	16.77	11.38	9.27	6.42	8.93	4.45	11.48	11.6	13.87										
PLACE1002721	75.97	53.95	40.28	41.2	67.12	33.39	16.52	34.39	33.55										
PLACE1002722	22.13	17.93	14.21	5.69	15.16	8.22	22.92	15.1	22.19										
PLACE1002726	7.85	7.95	6.75	3.03	4.47	1.65	5.1	2.92	3.72	**	**	-	-	-	-	-	-	-	-
PLACE1002756	11.64	12.77	9.75	6.41	12.37	7.42	9.89	7.27	9.88										
PLACE1002788	14.14	13.66	17.41	6.98	8.91	8.35	8.23	12.98	13.43	**	*	-	-	-	-	-	-	-	-
PLACE1002772	11.58	12.05	13.28	5.34	9.18	5.42	7.04	8.7	6.74	*	*	-	-	-	-	-	-	-	-
PLACE1002775	3.87	3.22	3.62	4.05	3.19	0.88	1.6	5	2.48										
PLACE1002780	8.73	8.1	8.81	2.91	6.16	2.82	3.16	6.37	4.26	*	*	-	-	-	-	-	-	-	-
PLACE1002782	15.09	18.6	18.22	6.13	9.37	3.85	11.71	13.03	12.96	**	*	-	-	-	-	-	-	-	-
PLACE1002784	41.39	43.8	44.4	31.78	35.79	20.44	25	22.44	21	*	*	-	-	-	-	-	-	-	-
PLACE1002785	26.08	25.22	25.67	11.64	12.89	13.19	17.32	17.1	14.49	**	**	-	-	-	-	-	-	-	-
PLACE1002811	45.78	45.83	45.46	25.72	32.66	23.15	17.78	22.39	22.68	**	**	-	-	-	-	-	-	-	-
PLACE1002815	15.82	13.42	19.22	7.58	7.46	5.89	11.61	7.34	8.29	**	*	-	-	-	-	-	-	-	-
PLACE1002816	4.88	2.68	5.34	6.88	10.83	8.75	3.44	3.56	2.71	**	+	-	-	-	-	-	-	-	-
PLACE1002822	12.2	11.93	11.07	6.15	11.75	10.16	7.41	12.24	8.12										
PLACE1002833	9.14	11.66	12.33	5.57	7.46	7.07	4.75	8.35	4.08	*	*	-	-	-	-	-	-	-	-
PLACE1002834	6.07	5.81	8.84	3.09	3.53	4.27	1.8	2.98	2.41	*	*	-	-	-	-	-	-	-	-
PLACE1002835	58.22	48.91	69.68	49.95	82.63	44.56	42.91	35.34	35.49	*	*	-	-	-	-	-	-	-	-
PLACE1002839	20.38	22.46	21.81	8.31	11	7.79	15.07	14.42	13.25	**	**	-	-	-	-	-	-	-	-
PLACE1002851	24.75	15.11	15.09	7.91	14.09	8.63	20.69	22.56	25.92										
PLACE1002853	17.64	15.34	22.8	5.62	8.45	8.1	5.28	8.18	4.17	**	**	-	-	-	-	-	-	-	-
PLACE1002881	24.88	14.92	26.75	13.38	16.28	13.04	14.2	20.46	17.04										
PLACE1002901	14.25	14.08	17.74	6.8	8.98	8.68	8.78	4.53	4.12	**	**	-	-	-	-	-	-	-	-
PLACE1002904	18.88	26.12	22.33	14.53	22.41	12.83	9.21	13.95	10.64	*	*	-	-	-	-	-	-	-	-
PLACE1002905	4.41	8.93	12	2.54	8.87	4.07	3.89	5.83	4.18										
PLACE1002908	13.22	18.1	16.63	15.4	12.99	11.14	8.47	5.63	8.35	**	*	-	-	-	-	-	-	-	-
PLACE1002911	10.97	9.2	9.02	8.7	11.96	7.89	5.04	4.71	5.5	**	*	-	-	-	-	-	-	-	-
PLACE1002941	15.88	10.4	17.73	6.14	8.22	5.77	14.58	18	12.89	*	*	-	-	-	-	-	-	-	-
PLACE1002980	23.85	20.5	17.79	16.86	17.44	5.49	24.31	14.61	17.93										
PLACE1002955	5	5.58	7.87	3.29	6.71	3.61	3.06	3.96	2.19	*	*	-	-	-	-	-	-	-	-
PLACE1002958	88.1	92.88	103.58	90.74	91.81	78.41	59.31	52.18	47.63	**	*	-	-	-	-	-	-	-	-
PLACE1002982	3.13	4.33	9.15	0.9	4.01	2.47	0.8	0.76	1.87										
PLACE1002987	35.4	50.14	51.41	21.55	34.46	20.97	31.98	28.21	37.19										
PLACE1002988	4.93	4.01	6.65	3.18	3.05	1.83	2.74	2.46	2.95	*	*	-	-	-	-	-	-	-	-
PLACE1002976	59.32	76.52	110.87	52.57	103.04	77.15	42.49	35.2	48.8										
PLACE1002991	6.98	4.41	7.67	2.11	5.68	3.18	2.77	5.13	6.68										
PLACE1002993	13.31	16.88	19.44	8.64	17.73	10.58	9.53	20.73	7.89										
PLACE1002995	33.54	26.83	25.33	24.84	36.88	25.62	15.6	28.74	20.69										
PLACE1003010	5.87	5.37	8.7	4.68	3.28	4.68	3.8	4.41	3.12										
PLACE1003025	19.13	20.65	29.6	10.42	22.81	11.62	14.92	22.04	12.57										
PLACE1003027	4.81	4.57	8.94	2.3	4.47	0.55	3.14	5.14	2.31										
PLACE1003044	13.45	13	13.89	6.33	7.16	6.74	3.17	4.52	2.59	**	**	-	-	-	-	-	-	-	-
PLACE1003045	18.2	13.12	16.27	11.21	14.83	10.08	8.58	7.68	5.73	**	*	-	-	-	-	-	-	-	-
PLACE1003052	16.26	13.65	16.48	5.57	12.75	6.74	15.39	12.34	13.31	*	*	-	-	-	-	-	-	-	-
PLACE1003083	29.68	32.06	29.45	16.75	26.42	15.92	23.57	20.32	20.14										
PLACE1003085	11.46	7.5	13.4	5.78	9.38	4.65	6.65	6.68	5.67										
PLACE1003092	13.44	11.78	13.5	8.16	11.43	7.08	8.07	7.87	6.9	*	*	-	-	-	-	-	-	-	-
PLACE1003097	6.17	6.46	16.32	3.05	5.89	3.08	3.13	5.88	4.77										
PLACE1003100	38.23	40.39	38.55	32.04	32.87	28.75	28.62	23.05	18.02	*	**	-	-	-	-	-	-	-	-
PLACE1003188	11.34	8.75	20.83	7.36	13.72	5.16	3.96	4.75	4.82										
PLACE1003115	7.34	6.75	7.41	3.32	6.32	2.75	1.85	0.73	1.92	**	*	-	-	-	-	-	-	-	-
PLACE1003120	12.49	11.78	16.03	7.83	8.21	4.78	11.87	11.32	7.77	*	*	-	-	-	-	-	-	-	-
PLACE1003185	7.09	5.05	7.89	3.8	2.81	2.79	2.84	3.42	1.44	*	*	-	-	-	-	-	-	-	-
PLACE1003136	68.81	77.15	81.78	65.88	72.38	60.39	44.74	39.52	47.1	**	*	-	-	-	-	-	-	-	-
PLACE1003141	9.02	4.98	11.12	4.59	4.78	5.17	4.62	5.32	3.84										

120 / 157

2679

2680

PLACE1003145	12.65	9.94	12.43	3.74	5.65	4.53	5.94	8.05	4.35	**	*	-	-
PLACE1003147	26.48	26.41	26.84	11.9	16.38	12.11	16.22	18.19	14.71	**	**	-	-
PLACE1003153	10.51	10.55	11.78	2.87	8.22	4.35	5.52	7.01	3.61	**	**	-	-
PLACE1003163	8.08	7.83	4.23	1.17	5.16	2.64	1.92	2.39	2.29	*	*	-	-
PLACE1003172	11.82	10	15.87	8.77	13.23	9.45	7.97	8.79	7.48	*	*	-	-
PLACE1003174	18.37	9.05	12.28	9.12	4.76	2.9	4.07	4.08	4.97	*	*	-	-
PLACE1003176	5.18	3.89	5.16	2.81	3.31	2.68	2.88	3.04	2.06	*	*	-	-
PLACE1003181	9.36	6.29	8.52	1.78	2.48	3.42	2.28	1.97	1.19	**	**	-	-
PLACE1003184	23.08	18.89	20.23	9.3	11.84	12.17	13.78	13.13	10.83	**	**	-	-
PLACE1003190	8.27	8.79	9.4	3.25	4.54	3.59	4.27	2.82	4.68	**	**	-	-
PLACE1003200	9.79	7.56	9.39	2.41	8.08	2.83	2.42	1.72	3.3	*	*	-	-
PLACE1003205	126.72	130.28	137.95	113.01	122.92	109.79	99.18	88	89.92	*	*	-	-
PLACE1003208	2.18	18.18	2.48	0.47	10.18	0.97	2.71	3.38	3.16	*	*	-	-
PLACE1003214	22.58	19.76	20.17	11.54	15.74	15.47	11.96	12.4	14.5	*	**	-	-
PLACE1003220	11.83	11.81	10.65	4.92	4.84	3.94	7.34	5.53	5.38	**	**	-	-
PLACE1003238	8.87	6.17	6.07	3.06	1.84	4.56	1.99	2.39	2.01	*	**	-	-
PLACE1003249	64.47	55.31	48	33.84	42.61	31.3	33.47	28.86	18.94	*	*	-	-
PLACE1003258	18.3	25.49	17.28	10.97	13.38	18.51	17.3	11.04	4.08	*	*	-	-
PLACE1003258	49.07	49.35	44.47	34.48	50.32	29.75	23.51	18.19	23.43	**	*	-	-
PLACE1003270	10.3	14.58	7.71	5.42	10.68	4.9	6.28	3.11	11.78	*	*	-	-
PLACE1003284	21.78	18.35	23.18	8.69	11.5	5.25	12.82	8.7	9.35	*	*	-	-
PLACE1003288	42.17	40.86	35.97	38.78	41.18	32.25	23.79	17.48	17.34	**	*	-	-
PLACE1003297	48.69	60.94	40.2	15.98	31.29	21.98	19.88	11.27	24.02	*	**	-	-
PLACE1003302	25.32	31.25	29.05	9.93	18.98	12.55	24.23	16.16	20.93	**	*	-	-
PLACE1003314	185.2	162.92	141.49	148.89	164.17	125.41	110.84	69.78	76.19	*	*	-	-
PLACE1003337	48.98	43.27	41.6	31.2	33.91	22.01	17.95	11.14	16.35	*	**	-	-
PLACE1003342	33.19	37.72	34.03	17.42	45.69	20.1	23.58	17.03	23.28	**	*	-	-
PLACE1003343	8.84	5.22	10.11	7.16	4.99	3.86	5.48	2.81	3.26	*	*	-	-
PLACE1003344	11.02	7.76	2.89	1.89	8.29	3.43	7.57	2.34	3.76	*	*	-	-
PLACE1003353	18.54	21.69	17.69	8.25	8.74	5.2	10.78	10.8	18.5	**	**	-	-
PLACE1003361	44.83	37.64	33.66	33.83	37.42	32.13	26.59	20.73	23.72	*	*	-	-
PLACE1003366	12	18.42	15.82	7.6	12.12	4.37	10.38	5.92	5.45	*	*	-	-
PLACE1003368	4.81	3.43	3.83	0.94	3.73	1.31	1.63	1.07	0.84	**	*	-	-
PLACE1003372	13.64	7.8	10.54	3.88	10.47	3.35	4.52	1.77	3.52	*	*	-	-
PLACE1003373	8.88	8.18	9.38	3.94	3.78	0.99	4.29	2.31	3.38	**	**	-	-
PLACE1003375	22.81	23.46	18.65	11.55	17.67	11.28	29.27	13.77	19.27	**	*	-	-
PLACE1003378	1.92	0.43	0.47	1.45	6.88	1.22	0.77	0.97	0.76	*	*	-	-
PLACE1003383	15.75	12.58	10.95	7.3	8.49	3.81	2.43	7.48	2.15	*	*	-	-
PLACE1003394	18.48	13.29	10.26	6.3	12.68	5.09	11.88	10.8	17.14	*	*	-	-
PLACE1003401	8.87	8	5.04	3.86	4.75	2.93	5.66	3.3	5.47	*	*	-	-
PLACE1003405	7.8	8.07	5.81	3.39	5.27	4.58	4.12	1.47	2.3	*	*	-	-
PLACE1003407	10.24	7.49	7.79	2.43	3.28	2.12	4.16	2.37	3.6	**	**	-	-
PLACE1003420	7.71	5.57	6.39	3.78	6.36	2.21	2.89	1.88	2.26	**	*	-	-
PLACE1003428	6.55	5.9	4.45	2.19	4.16	0.9	3.49	2.31	8.87	*	*	-	-
PLACE1003432	51.6	69.55	75.77	36.78	44.92	44.22	42.27	64.14	52.55	*	*	-	-
PLACE1003438	1.98	2.35	1.48	2.21	4.73	1.43	4.08	0.93	0.7	*	*	-	-
PLACE1003452	3.01	3.41	4.3	1.8	3.84	2.08	2.34	2.25	1.84	*	*	-	-
PLACE1003464	3.87	4.87	5.58	2.64	4.5	3.92	2.57	2.43	1.33	*	*	-	-
PLACE1003465	2.82	2.41	4.04	1.6	1.9	1.64	1.76	0.8	1.88	*	*	-	-
PLACE1003466	2.38	4.17	3.83	2.08	10.37	2.22	2.4	1.94	0.64	*	*	-	-
PLACE1003467	4.78	5.09	4.78	4.08	6	2.64	6.38	3.48	2.92	*	*	-	-
PLACE1003478	8.15	2.11	2.68	1.92	3.98	2	5.39	1.91	4.04	*	*	-	-
PLACE1003484	8.84	8.17	8.98	4.38	6.73	3.1	4.12	4.68	4.63	**	*	-	-
PLACE1003493	83.14	41.9	88.74	60.01	72.45	45.07	64.55	71.53	53.96	*	*	-	-
PLACE1003503	7.38	5.03	6.29	1.12	3.91	3.51	2.38	3.64	2.4	*	*	-	-
PLACE1003505	32.2	45.22	38.51	20.24	38.17	23.68	24.44	34.5	27.01	*	*	-	-
PLACE1003516	7.02	7.8	7.71	2.88	5.85	3.76	2.24	2.88	3.88	*	**	-	-
PLACE1003518	4.08	1.81	6.55	3.83	3.85	1.91	2.58	1.64	0.34	*	*	-	-
PLACE1003520	12.72	16.98	9.15	8.71	16.58	8.45	4.85	6.39	5.22	*	*	-	-
PLACE1003521	14.95	10.13	14.17	8.02	7.53	5.58	6.77	8.38	4.75	*	*	-	-
PLACE1003526	20.17	19.13	25.59	15.85	14.55	12.63	20.97	18.71	22.38	*	*	-	-
PLACE1003528	35.65	25.12	43.38	21.83	30.38	28.64	18.82	24.64	18.65	*	*	-	-
PLACE1003529	11.19	18.69	21.6	20.8	18.68	31.76	18.24	15.7	15.44	*	*	-	-
PLACE1003537	3.16	4.02	8.21	3.24	3.64	3.89	1.78	2.66	2.32	*	*	-	-

121 / 157

2681

2682

PLACE1003549	16.21	19.62	27.74	11.72	18.71	13.09	12.65	15.02	13.74				
PLACE1003552	12.63	20.69	18.89	10.17	18.28	8.51	8.12	8.44	8.9	*	-		
PLACE1003566	8.49	12.35	9.77	7.53	10.49	7.08	3.77	4.68	6.9	*	-		
PLACE1003568	5.3	4.9	19.78	3.8	7.59	3.87	5.3	6.3	4.21				
PLACE1003573	32	27.25	37.46	17.52	23.28	16.23	15	18.28	19.13	*	**	-	-
PLACE1003575	66.07	60.03	70.99	42.8	62.9	38.6	75.17	59.46	62.58				
PLACE1003583	8.33	4.65	6.88	3.47	6.26	5.07	2.38	3.4	4.1	*	-		
PLACE1003584	12.09	10.68	18.02	7.17	8.48	19.2	8.54	7.15	2.79				
PLACE1003592	29.77	33.55	28.89	21.34	26.31	15.85	12.59	8.49	12.12	*	**	-	-
PLACE1003593	7.15	3.54	8.27	3.34	2.24	1.83	3.11	5.19	5.45				
PLACE1003594	7.18	8.49	9.31	5.03	6.61	3.38	4.99	3.18	2.37	*	**	-	-
PLACE1003596	5.11	3.13	4.25	2.82	5	3.14	3.75	4.1	3.78				
PLACE1003598	31.57	28.75	31.55	14.81	20.8	17.31	5.31	17.81	7.34	**	**	-	-
PLACE1003602	6.51	2.47	7.47	1.84	4.91	0.98	2.4	4.6	2.56				
PLACE1003605	10	8.14	7.29	5.28	8.85	3.75	6.45	5.07	2.93				
PLACE1003611	15.94	16.54	24.25	11	9.1	12.46	10.14	10.62	12.83	*	-		
PLACE1003618	7.78	7.74	12.1	3.49	8.12	2.68	4.58	5.29	8.93	*	-		
PLACE1003625	14.19	21.53	18.27	11.53	22.81	17.82	13.8	9.54	11.66				
PLACE1003626	11.55	14.38	14.6	5.94	11.97	7.26	4.67	4.49	2.94	**	-		
PLACE1003630	3.66	4.65	3	4.82	4.58	3.02	3.68	3.68	1.88				
PLACE1003635	8.33	7.29	10.19	5.35	5.59	4.6	3.51	4.48	4.17	*	**	-	-
PLACE1003638	4.33	3.09	4.83	3.88	8.02	2.88	2.01	3.6	1.35				
PLACE1003644	36.03	24.77	24.25	19.14	20.11	17.96	8.77	10.6	6.97	**	-		
PLACE1003654	75.56	74.67	74.52	77.81	72.17	67	35.39	33.92	31.08	**	-		
PLACE1003656	7.81	5.19	8.37	3.08	4.32	3.8	1.18	1.63	2.96	*	-		
PLACE1003660	16.26	21.8	21.11	12.32	17.1	13.06	14.44	12.72	12.99	*	-		
PLACE1003669	12.05	8.79	12.65	4.93	8.33	7.48	3.9	4.68	4.41	**	-		
PLACE1003670	110.16	83.05	76.99	55.55	62.05	50.36	48.6	63.39	52.6	*	-		
PLACE1003671	11.47	3.49	4.69	7.13	4.59	4.35	3.71	2.69	2.22				
PLACE1003697	8.65	10.83	16.05	9.21	10.08	8.26	6.51	4.97	8.75	*	-		
PLACE1003704	10.59	7.92	10.14	6.15	6.81	6.61	4.22	3.63	3.88	*	**	-	-
PLACE1003709	9.06	8.99	10.78	5.34	8.7	5.47	3.53	3.38	1.83	**	**	-	-
PLACE1003711	28.05	31.28	32.33	21.12	31.77	27.07	16.12	15.79	14.21	**	-		
PLACE1003723	27.32	34.97	28.91	18.16	26.8	21.24	17.63	17.4	10.81	**	-		
PLACE1003724	29.33	26.12	25.65	19.94	34.28	21.71	20.68	13.83	15.71	**	-		
PLACE1003737	8.24	6.31	9.31	2.69	4.06	2.72	3.31	2.56	3.28	**	**	-	-
PLACE1003738	5.02	4.71	5.03	6.78	2.78	9.43	1.8	2.2	2.93	**	-		
PLACE1003742	17.85	18.69	20.82	14.24	7.57	8.14	10.49	8.98	8.16	*	**	-	-
PLACE1003744	5.48	2.82	3.4	0.68	1.04	1.08	1.74	1.94	0.38	*	-		
PLACE1003758	23.32	22.73	21.79	9.9	15.26	16.38	15.29	14.75	13.36	*	**	-	-
PLACE1003760	40.1	57.96	44.54	28.85	37.15	22.14	11.09	14.54	11.41	**	-		
PLACE1003762	85.35	58.09	77.88	35.3	88.27	46.92	46.48	32.75	38.3	*	-		
PLACE1003765	93.01	107	70.69	55.75	97.14	46.35	33.33	44.53	44.51	*	-		
PLACE1003768	34.07	21.23	23.34	10.87	13.6	6.67	25.2	17.48	28.3	*	-		
PLACE1003771	5.38	5.57	5.27	3.47	7.27	2.71	1.77	1.77	2.44	**	-		
PLACE1003772	5.63	7.96	4.89	1.17	7.74	2.12	3.12	2.24	2.28	*	-		
PLACE1003783	5.15	3.84	3.95	2.38	3.83	1.92	1.65	1.17	0.84	**	-		
PLACE1003784	5.69	6.98	6.68	2.81	4.01	2.73	3.1	2.42	1.88	**	**	-	-
PLACE1003788	34.98	33.59	27.31	18.49	15.1	21.27	12.2	5.92	10.73	*	**	-	-
PLACE1003795	26.48	33.02	23.72	12.5	28.84	12.28	23.33	16.94	18.45				
PLACE1003827	9.74	8.51	10.55	21.41	16.48	10.33	5.7	3.14	1.91	**	-		
PLACE1003833	14.14	10.78	12.27	5.74	8.62	5.65	5.59	6.29	7.37	*	**	-	-
PLACE1003839	259.31	241.18	222.89	217.52	173.45	205.17	183.55	138.4	151.61	**	-		
PLACE1003845	69.72	60.2	118.04	40.89	28.14	26.89	48.08	43.86	45.73				
PLACE1003850	9.53	11.45	9.42	3.3	8.43	5.88	6.51	3.87	5.97	*	-		
PLACE1003852	32.49	28.32	26.73	21.78	30.69	26.63	18.45	14.12	17.92	**	-		
PLACE1003858	19.9	13.83	13.88	8.03	12.82	5.24	7.08	4.19	7.71	*	-		
PLACE1003861	40.22	66.3	33.51	12.47	57.37	18.72	33.23	18.67	19.13				
PLACE1003864	34.51	84.78	122.59	61.83	70.16	56.03	69.32	56.31	42.37				
PLACE1003870	9.43	8.5	5.9	2.94	3.32	3.4	5.69	3.88	5.74	*	-		
PLACE1003885	10.07	6.78	3.91	3.5	6.22	3.91	3.49	3.72	6.33				
PLACE1003886	12.68	12.78	10.57	6.28	9.54	3.51	13.88	8.88	12.28	*	-		
PLACE1003888	15.48	13.6	17.45	7.94	12.36	6.23	14.56	8.73	9.07	*	-		
PLACE1003892	6.59	4.85	8.21	5.64	7.14	2.75	3.77	2.15	4.24				

122 / 157

2683

PLACE1003900	9.05	6.59	6.53	3.41	9.33	4.49	4.41	6.87	4.79				
PLACE1003902	9.43	6.06	7.46	2.35	3.67	1.78	3.96	1.76	3.05	*	*	-	-
PLACE1003903	17.33	10.38	14.55	9.42	10.87	10.03	9.08	6.58	7.77	**	**	-	-
PLACE1003915	14.67	18	11.19	12.84	10.38	9.98	8.5	5.29	8.8	*	*	-	-
PLACE1003918	10.58	6.82	8.73	7.34	5.29	3.45	7.05	5.69	4.94				
PLACE1003923	11.24	8.95	4.47	2.95	4.6	4.65	4.65	3.26	3.87				
PLACE1003932	9.02	5.88	4.39	2.18	9.39	4.8	8.3	4.31	5.14				
PLACE1003938	11.92	7.46	8.85	2.47	4.23	4.38	9.51	9.11	10.65	*	*	-	-
PLACE1003968	11.6	10.3	9.01	5.2	6.1	4.37	6.77	5.83	6.22	**	**	-	-
PLACE1003968	4.88	5.07	4.71	0.68	3.26	0.8	3.51	1.87	0.72	*	*	-	-
PLACE1004018	4.89	2.45	2.55	4.7	8.1	2.18	2.99	1.85	1.89				
PLACE1004020	12.24	14.87	17.88	7.54	8.19	5.65	7.35	9.99	10.06	*	*	-	-
PLACE1004028	2.49	1.88	2.35	1.78	7.14	2.28	1.01	1.49	0.82	*	*	-	-
PLACE1004034	6.5	4.91	7.64	2.7	5.45	2.86	3.71	3.07	1.75	*	*	-	-
PLACE1004042	5.94	2.98	4.63	2.04	4.08	3.55	3.58	3.9	2.95				
PLACE1004078	6.26	4	5.13	2.35	3.67	1.82	1.94	3.68	4.31	*	*	-	-
PLACE1004103	4.18	6.73	8.68	2.05	3.45	2.36	1.89	0.91	3.01	*	*	-	-
PLACE1004104	0.6	1.12	3	1.89	7.2	1.01	1.88	2.79	1.52				
PLACE1004113	0.85	1	0.69	0.62	4.18	1	1.54	0.77	1.59				
PLACE1004114	12.27	11.63	12.84	9.04	6.13	6.15	17.17	8.3	11.59	**	**	-	-
PLACE1004118	25.58	29.6	34.14	10.89	22.13	12.84	16.2	18.15	18.4	*	*	-	-
PLACE1004128	6.77	8.44	8.16	3.47	7.23	7.78	2.81	3.08	3.14	**	**	-	-
PLACE1004130	9.38	4.88	10.86	8.2	7.53	9.87	4.28	6.71	6.1				
PLACE1004149	2.51	2.43	6.21	2.78	5.35	2.6	3.3	3.26	2.01				
PLACE1004166	13.06	9.08	13.29	7.39	11.05	6.74	9.05	8.4	10.62				
PLACE1004180	20.97	18.94	23.27	14.03	30.3	7.33	16.87	16.65	14.66	*	*	-	-
PLACE1004181	7.06	8.18	8.05	5.83	7.98	4.77	4.77	4.44	4.89	**	**	-	-
PLACE1004188	41.1	38.11	42.08	18.36	22.19	21.87	24.58	28.32	30.02	**	**	-	-
PLACE1004188	32.92	30.07	31.55	16.2	15.66	13.35	21.23	25.32	27.11	**	*	-	-
PLACE1004170	14.53	21.92	19.94	10.32	12.85	10.61	11.03	18.31	8.2	*	*	-	-
PLACE1004178	2.75	3.62	3.97	2.55	6.19	3.35	0.34	1.55	0.72	**	**	-	-
PLACE1004183	18.95	43.53	45.99	27.78	51.64	25.82	21.25	25.43	23.82	**	**	-	-
PLACE1004197	5.75	5.24	7.84	2.14	4.97	3.74	3.04	2.49	3.28	*	*	-	-
PLACE1004199	21.75	27.66	12.12	12.42	24.13	11.64	10.47	10.88	12.46				
PLACE1004203	11	13.62	10.81	4.23	10.81	6.68	10.74	11.32	12.32				
PLACE1004242	20.49	18.48	21.11	6.91	6.93	18.71	6.23	11.27	9.24	*	**	-	-
PLACE1004249	6.73	11	11.25	4.47	10.28	6.2	5.79	8.22	6.89	*	*	-	-
PLACE1004255	32.88	15.95	34.9	17.52	26.48	21.84	27.95	27.86	29.02	*	*	-	-
PLACE1004258	12.54	10.51	15.03	5.99	6.51	6.23	3.84	5.04	3.23	**	**	-	-
PLACE1004257	3.25	14.27	11.31	3.38	8.12	8.15	2.7	9.49	4.03				
PLACE1004258	11.89	28.37	21.62	8.78	18.18	12.25	12.68	13.44	9.65				
PLACE1004270	8.17	8.34	11.88	6.47	5.77	11.76	8.27	3.74	6.39				
PLACE1004272	1.52	1.56	3.17	1.04	3.51	0.48	1.56	1.06	1.78				
PLACE1004273	23.31	38.14	20.67	17.38	14.34	7.01	17.2	17.07	20.56				
PLACE1004274	12.13	13.19	15.05	6.12	12.19	6.04	6.47	11.78	6.75				
PLACE1004277	16.65	17.3	18.15	12.2	21.29	13	13.46	15.12	20.1				
PLACE1004279	20.99	17.87	19.27	8.98	30.05	13	8.21	11.3	11.18	**	**	-	-
PLACE1004282	12.55	12.53	18.68	4.95	9.12	6.25	8.7	8.14	5.89	*	**	-	-
PLACE1004284	9.58	14.66	15.53	8.95	7.35	5.65	6.2	8.82	3.1	*	*	-	-
PLACE1004289	11.23	10.79	11.74	2.16	5.71	3.35	2.35	6.14	2.14	**	**	-	-
PLACE1004299	19.52	18.15	18.38	8.87	22.44	9.43	11.73	9.62	8.57	**	**	-	-
PLACE1004302	6.5	4.57	5.26	4.18	7	3.55	1.58	4.16	0.55	*	*	-	-
PLACE1004305	7.18	4.88	7.78	4.02	11.98	4.38	3.44	4.82	2.94	*	*	-	-
PLACE1004316	21.08	20.32	25.43	14.83	15.07	12.7	13.59	13.17	12.8	*	**	-	-
PLACE1004322	8.54	3.44	10.02	13.05	22.34	1.38	2.62	7.35	3.43				
PLACE1004325	13.79	15.82	21.16	8.25	8.41	6.63	10.63	11.11	9.78	*	*	-	-
PLACE1004332	13.76	26.81	27.83	10.52	16.87	4.88	9.28	8.37	7.2	*	*	-	-
PLACE1004336	8.25	8.04	10.31	3.17	5.77	3.05	2.53	4.29	2.91	**	**	-	-
PLACE1004346	11.14	15.86	16.19	8.6	6.7	8.17	14.67	2.81	3.65	*	*	-	-
PLACE1004358	4.9	6.8	5.9	4.68	3.48	2.52	2.87	2.07	1.85	**	**	-	-
PLACE1004376	4.37	7.68	6.28	3.82	16.23	3.24	18.48	3.42	3.06				
PLACE1004384	20.82	15.88	20.1	10.2	14.54	11.78	10.47	11.55	9.77	*	**	-	-
PLACE1004385	18.21	25.32	28.3	8.24	11.85	7.12	18.49	9.57	9.87	*	*	-	-
PLACE1004388	13.03	15.84	14.98	8.25	11.49	8.38	7.85	6.97	4.61	*	**	-	-

2684

2685

2686

PLACE1004405	6.83	7.82	8.42	1.74	2.87	2.17	1.79	1.77	1.06	00	00	-	-
PLACE1004407	11.75	11.43	14.15	6.35	10.59	6.28	6.13	6.07	6.47	*	00	-	-
PLACE1004424	6.94	8.4	8.66	2.02	6.19	2.69	3.43	3.43	2.34	*	00	-	-
PLACE1004425	15.08	13.72	13.15	6.65	11.37	7.31	11.7	10.39	11.17	*	0	-	-
PLACE1004427	9.79	7.13	8.53	5.79	4.82	4.58	2.31	1.96	2.72	*	00	-	-
PLACE1004428	8.46	5.33	5.57	3.25	3.29	3.21	2.85	2.65	2.1	*	0	-	-
PLACE1004433	6.57	6.12	7.26	2.15	3.01	2.68	2.76	2.36	3.32	*	00	-	-
PLACE1004435	5.31	5.07	8.8	2.78	3.41	3.61	2.83	0.69	1.36	*	0	-	-
PLACE1004437	123.68	158.84	144.68	118.58	167.95	88.95	49.84	58.42	68.77	00	00	-	-
PLACE1004441	10.02	12.73	11.55	5.63	11.08	5.24	5.14	4.58	2.83	00	00	-	-
PLACE1004446	4.95	6.73	4.89	2.19	3.83	1.62	2.79	2.51	2.66	*	00	-	-
PLACE1004450	17.21	13.77	11.35	7.43	9.82	10.35	13.72	15.81	16.38	00	00	-	-
PLACE1004451	21.93	21.83	22.42	14.24	15.13	14.34	10.39	9.42	10.97	00	00	-	-
PLACE1004456	5.68	3.89	4.79	2.22	5.14	3.7	4.95	1.66	2.31	0	0	-	-
PLACE1004458	6.57	4.16	7.77	2.1	5.76	2.18	2.41	3.03	1.64	0	0	-	-
PLACE1004460	6.93	5.13	4.79	2.01	4.05	1.65	3.51	2.54	1.12	0	0	-	-
PLACE1004467	10.11	11.82	9.53	2.74	4.49	5.1	3.12	1.44	1.55	00	00	-	-
PLACE1004471	14.09	22.14	14.79	8.38	18.33	11.21	5.81	8.78	5.83	0	0	-	-
PLACE1004473	8.27	14.31	9.68	3.42	8.92	3.77	4.46	5.47	8.39	0	0	-	-
PLACE1004475	4.2	2.67	2.47	2.6	8.42	6.68	3.42	2.58	2.7	0	0	-	-
PLACE1004482	42.48	39.9	32.17	18.98	31.24	19.51	41.1	31.68	45.4	0	0	-	-
PLACE1004481	3.4	5.37	4.92	2.26	2.87	1.21	3.11	1.77	1.09	0	0	-	-
PLACE1004482	14.75	10.5	4.25	8.39	18.03	7.27	8.28	2.19	2.31	0	0	-	-
PLACE1004506	18.11	19.82	21.23	12.46	23.49	16.62	17.33	12.61	13.21	0	0	-	-
PLACE1004507	14.62	14.01	19.19	12.21	20.75	13.12	6.26	8.93	6.09	0	00	-	-
PLACE1004510	8.27	7.11	7.77	4.2	4.81	1.36	5.06	2.94	3.49	0	00	-	-
PLACE1004516	5.75	6.87	3.83	4.51	8.48	2.72	4.89	2.56	3.06	0	0	-	-
PLACE1004518	9.64	6.39	6.6	5.58	7.29	2.45	2.45	4.03	4.48	0	0	-	-
PLACE1004519	45.05	38.06	29.43	18.22	21.37	17.38	37.02	28.03	42.7	0	0	-	-
PLACE1004520	6.65	5.88	4.83	5	23.3	3.26	7.89	2.48	5.32	00	00	-	-
PLACE1004530	8.66	8.01	8.19	4.33	4.87	3.07	5.74	4.33	5.72	00	00	-	-
PLACE1004545	16.35	9.42	10.89	5.28	8.16	4.82	7.69	6.29	6.88	0	0	-	-
PLACE1004547	8.88	4.99	7.28	3.79	10.89	2.44	2.07	2.03	2.03	0	0	-	-
PLACE1004548	8.06	7.34	7.17	1.99	3.55	1.53	3.36	1.74	4.09	00	00	-	-
PLACE1004550	9.29	4.74	4.67	6.39	7.24	2.64	11.12	8.11	5.12	0	0	-	-
PLACE1004551	28.28	24.92	17.22	13.36	20.97	8.68	22.61	16.23	21.2	0	0	-	-
PLACE1004559	38.54	25.35	19.12	16.44	15.14	18.5	25.43	24.95	33.62	0	0	-	-
PLACE1004582	50.28	46.32	50.21	54.16	58.41	59.84	35.71	42.24	45.43	0	0	-	-
PLACE1004584	12.01	9.86	9.29	4.62	9.04	5.79	9.27	6.54	10.21	00	00	-	-
PLACE1004604	17.66	17.86	15.11	9.74	9.81	7.68	16.75	11.47	16.51	00	00	-	-
PLACE1004611	6.51	3.84	4.64	1.34	7.9	1.38	3.77	0.9	3.89	0	0	-	-
PLACE1004629	7.22	4.23	7.21	2.88	5.64	1.27	4	2.37	6.88	0	0	-	-
PLACE1004630	5.71	5.71	5.05	2.41	3.38	0.92	6.59	3.07	5.19	0	0	-	-
PLACE1004637	16.22	15.19	15.73	4.5	8.73	18.68	5.83	11.18	5.64	0	0	-	-
PLACE1004645	16.19	10.08	17.1	10.38	11.57	13.97	10.9	12.36	6.68	00	00	-	-
PLACE1004646	4.97	5.84	4.39	4.13	5.71	3.39	2.58	3.34	2.67	00	00	-	-
PLACE1004648	5.49	4.11	3.98	3.64	5.18	4.29	1.91	1.5	0.14	00	00	-	-
PLACE1004655	3.31	1.38	2.13	1.15	3.39	0.72	0.74	1.08	1.47	00	00	-	-
PLACE1004658	61.59	67.15	67	38.54	49.97	38.25	52.62	54.12	65.08	00	00	-	-
PLACE1004664	1.18	4.36	2.93	2.88	5.73	2.7	1.85	2.13	1.51	00	00	-	-
PLACE1004672	28.88	30.42	26.6	16.98	20.16	13.59	16.28	11.87	13.82	00	00	-	-
PLACE1004674	4.27	1.5	4.11	2.41	3.15	1.89	1.56	2.98	3.87	0	0	-	-
PLACE1004681	8.54	6.41	13.05	7.27	21.72	12.91	4.84	4.9	3.36	0	0	-	-
PLACE1004686	4.34	3.3	4.81	2.4	3.38	2.38	2.29	3.83	0.39	0	0	-	-
PLACE1004690	6.4	6.12	8.6	4.32	13.98	8.68	2.73	4.58	2.78	0	0	-	-
PLACE1004691	12.44	14.52	17.78	7.94	13.42	7.13	8.04	12.14	11.56	00	00	-	-
PLACE1004693	8.4	10.27	9.88	5.82	5.16	5.28	2.45	3.25	2.73	00	00	-	-
PLACE1004701	144.88	188.18	97.68	180.88	187.91	84.32	88.33	73.95	139.9	00	00	-	-
PLACE1004705	3.47	3.01	2.93	2.39	3.98	1.27	1.92	3.49	3.87	00	00	-	-
PLACE1004708	48.88	43.13	45.03	31.98	28.65	23.68	15.08	18.44	11.77	00	00	-	-
PLACE1004718	20.04	14.83	24.39	8.19	12.11	8.27	8.37	8.78	9.57	0	0	-	-
PLACE1004722	5.57	7.39	10	4.45	8.54	3.93	3.67	3.69	3.48	0	0	-	-
PLACE1004736	16.71	17.15	24.67	12.34	13.91	6.89	9.19	22.16	13.53	0	0	-	-
PLACE1004737	5.22	7.74	7.78	3.83	6.24	1.98	3.08	3.38	3.06	0	0	-	-

124 / 157

2687

PLACE1004740	5.7	7.45	13.2	8.48	9.7	6.78	5.61	4.01	4.39										
PLACE1004743	25.2	29.5	32.11	15.32	29.52	15.01	26.22	29.41	28.91										
PLACE1004751	26.3	37.3	36.23	17.85	40.34	18.49	27.84	27.38	23.37										
PLACE1004757	27.21	29.89	35.01	20.07	11.76	13.22	19.63	14.54	15.78										
PLACE1004761	38.82	28.57	36.7	16.84	22.51	18.17	11.71	16.47	11.41										
PLACE1004773	6.34	3	4.53	3.18	3.86	2.2	2.15	2.65	2.95										
PLACE1004775	6.98	4.5	11.09	2.47	5.26	2.58	2.74	2.24	1.32										
PLACE1004777	22.12	29.48	28.12	13.38	15.93	13.39	17.54	29.58	18.88										
PLACE1004793	38.61	50.6	39.99	29.77	37.84	25.83	24.47	24.63	21.5										
PLACE1004796	8.45	9.48	8.63	3.38	9.79	3.2	3.16	2.68	3.41										
PLACE1004804	5.98	3.66	3.68	1.07	6.07	2.59	2.15	2.35	2.21										
PLACE1004813	7.54	8.6	8.99	4.2	5.66	4.51	4.87	4.35	4.23										
PLACE1004814	4.13	4.9	5.89	3.27	11.52	2.25	2.11	31.08	4.24										
PLACE1004815	5.57	3.05	6.31	3.48	4.55	1.54	4.88	2.03	1.26										
PLACE1004816	10.65	12.52	15.48	48.15	13.59	7.88	11.01	5.49	24.95										
PLACE1004824	10.1	10.18	14.83	3.84	7.0	4.37	6.87	4.52	8.59										
PLACE1004827	6.33	6.31	10.2	1.77	3.73	0.89	1.94	4.38	3.45										
PLACE1004838	18.51	24.59	14.94	5.88	11.83	5.72	3.63	5.6	4.83										
PLACE1004839	32.68	37.85	28.97	17.25	33.86	18.53	18.4	22.65	17.82										
PLACE1004840	26.87	33.84	24.99	11.9	21.32	18.63	17.68	19.77	20.11										
PLACE1004842	29.36	29.36	38.71	28.03	17.22	20.39	11.25	16.65	13.58										
PLACE1004850	12.98	12.93	18.16	9.69	18.5	6.77	8.83	9.34	8.77										
PLACE1004888	19.8	16.22	20.8	8.53	18.93	9.03	11.19	11.11	9.47										
PLACE1004885	7.93	7.22	6.94	1.5	3.78	3.6	1.94	2.3	1.73										
PLACE1004886	8.48	8.07	17.49	4.04	4.53	2.9	4.63	4.47	2.5										
PLACE1004887	11.79	13.7	11.52	8.38	18.2	7	3.69	4.3	5.52										
PLACE1004888	6.25	6.09	7.72	0.93	8.2	4.59	2.58	2.25	1.01										
PLACE1004900	11.28	8.08	8.89	8.58	8.42	4.7	4.98	5.45	3.82										
PLACE1004902	7.13	7.88	10.7	4.87	6.92	2.81	4.29	4.21	2.64										
PLACE1004904	23.08	23.38	37.33	15.52	22.71	12.92	16.1	22.74	25.02										
PLACE1004911	11.74	10.87	12.86	7.47	12.66	7.84	6.28	5.55	4.09										
PLACE1004918	12.82	12.15	14.37	4.58	7.86	4.67	7.24	7.07	6.68										
PLACE1004918	17.31	18.68	18.94	7.38	10.01	8.44	8.32	5.58	6.18										
PLACE1004930	6.79	6.08	5.8	1.3	2	3.65	2.19	1.6	1.33										
PLACE1004934	14.88	19.18	12.29	5.42	10.92	8.11	4.81	8.88	7.47										
PLACE1004937	18.93	13.85	13.4	11.81	10.98	11.51	15	13.14	16.45										
PLACE1004949	5	5.22	6.84	6.78	3.31	3.09	4.44	4.05	3.5										
PLACE1004969	13.91	13.85	11.65	11.71	12.45	9.25	4.82	9.25	5.58										
PLACE1004970	28.75	33.22	29.53	13.51	28.35	23.54	15.11	12.68	14.15										
PLACE1004972	13.28	11.27	10.14	5.85	6.98	5.82	4.89	3.42	1.64										
PLACE1004974	18.47	29.85	28.89	16.44	23.94	17.53	14.71	17.1	12.72										
PLACE1004975	12.92	11.68	12.08	5.89	13.8	6.88	5.54	3.55	3.43										
PLACE1004979	7.78	7.47	6.88	3.07	7.65	3.78	4.2	4.58	6.5										
PLACE1004982	30.58	23.61	26.29	14.3	18.23	11.42	16.82	24.16	20.51										
PLACE1004985	29.56	24.84	25.1	15.88	18.26	12.6	21.78	17.15	20.25										
PLACE1005003	16.43	14.5	16.78	6.69	10.88	11.47	1.79	5.8	8.41										
PLACE1005004	5.25	4.21	4.06	1.44	3.4	2.25	2.78	1.23	1.09										
PLACE1005005	17.79	13.88	18.79	11.99	10.57	8.86	7.49	5.3	5.99										
PLACE1005011	10.94	14.04	9.8	3.35	8.24	6.44	3.48	2.32	0.6										
PLACE1005028	31.74	35.88	28.88	14.82	24.65	14.87	35.74	25.33	28.57										
PLACE1005027	26.29	29.23	15.87	7.97	18.55	9.09	25.84	19.81	24										
PLACE1005031	38.12	16.05	14.24	9.18	25.55	17.12	7.22	9.47	10.38										
PLACE1005035	13.88	12.84	17.42	8.68	15.84	10.29	11.23	7.16	8.77										
PLACE1005041	111.13	87.68	72.07	36.83	51	43.47	99.34	93.97	97.87										
PLACE1005045	27.6	29	18.07	15.08	19.65	20.68	22.18	13.25	24										
PLACE1005047	60.84	34.08	35.58	25.13	37.56	21.77	26.01	15.91	25.77										
PLACE1005052	10.39	12.55	10.8	4.67	15.48	6.61	10.9	9.05	18										
PLACE1005055	24.31	33.8	18.61	16.75	10.79	11.1	19.54	11.48	10.17										
PLACE1005086	7.67	8.41	7	12.21	61.68	20.27	25.44	4.84	8.48										
PLACE1005077	8.62	6.97	5.62	4.44	4.98	3.2	5.42	5.71	7.8										
PLACE1005085	10.11	4.48	8.84	3.42	3.7	2.05	2.8	5.01	2.88										
PLACE1005086	11.97	11.74	11.67	7.31	18.85	7.84	8.74	6.83	9.82										
PLACE1005088	86.57	83.58	74.65	79.4	89.3	75.33	58.8	52.44	57.51										
PLACE1005089	8.31	4.13	4.27	1.8	3.13	4.97	2.18	0.55	2.04										

125 / 157

2688

2689

2690

PLACE1005101	11.5	9.29	11.94	9.18	4.8	5.47	5.22	3.69	6.78	*	**	-	-
PLACE1005102	22.17	25.88	27.54	8.07	7.69	6.94	7.15	5.65	7.82	**	**	-	-
PLACE1005108	10.15	7.73	6.02	2.99	6.05	3.33	4.8	7.23	6.47			-	-
PLACE1005110	3.97	2.39	3.45	3.59	3.70	1.51	2.14	1.33	2.15			-	-
PLACE1005111	23.94	21.69	16.98	14.04	21.5	8.57	20.32	16.74	23.71			-	-
PLACE1005123	16.94	14.46	12.95	5.45	9.5	5	6.89	4.29	5.82	*	**	-	-
PLACE1005124	9.14	6.62	8.15	2.08	5.47	1.8	6.61	3.48	5.33	*		-	-
PLACE1005128	6.83	4.81	4.87	1.09	3.42	1.91	3.3	3.83	2.82	*	*	-	-
PLACE1005130	12.43	17.68	12.68	4.58	11.05	5.79	12.55	5.44	10.87			-	-
PLACE1005141	6.07	3.99	6.18	8.23	2.88	1.1	9.65	2.17	2.82			-	-
PLACE1005146	10.52	9.7	6.88	8.24	10.69	5.88	5.25	8.14	5.49			-	-
PLACE1005152	91.15	76.64	75.79	51.08	95.85	52.59	42.4	38.1	38.75	**		-	-
PLACE1005157	6.11	3.75	6.08	2.82	4.88	2.4	1.9	2.79	1.7	*		-	-
PLACE1005162	24.79	23.25	27.03	17.82	28.77	20.32	10.52	15.51	11.65	**		-	-
PLACE1005170	9.55	7.54	6.99	11.08	19.85	14.4	3.82	8.2	5.94			-	-
PLACE1005176	5.55	5.17	6.85	2.52	4.13	3.01	2.09	2.54	1.28	*	**	-	-
PLACE1005181	10.56	18.07	15.09	7.48	9.08	8.01	10.91	12.85	10.11	*		-	-
PLACE1005184	3.64	4.4	5.5	5.28	11.77	4.28	3.84	3.69	3.06			-	-
PLACE1005185	5.98	5.07	5.66	2.88	2.54	3.28	3.84	4.33	4.72	**	*	-	-
PLACE1005187	20.63	14.71	18.55	8.5	10.68	10.85	12.23	12.87	7.61	*	*	-	-
PLACE1005189	8.28	7.72	11.92	4.24	7.19	3.99	6.84	7.52	5.99			-	-
PLACE1005193	19.26	44.11	39.59	31.34	38.98	28.91	38.03	40.58	31.59	*	+	-	-
PLACE1005200	1391.4	1317.2	1202.3	1770.3	2457.3	1744.3	1189.6	1397.9	588.38	*		-	-
PLACE1005205	48.27	41.57	88.17	48.38	52.86	50.82	27.19	13.53	58.32			-	-
PLACE1005216	10.1	9.74	10.26	6.79	10.41	5.95	3.4	3.26	4.05	**		-	-
PLACE1005223	36.88	44.94	37.06	23.02	26.39	16.59	23.41	25.05	19.76	*		-	-
PLACE1005225	5.34	5.81	6.83	4.25	3.02	5.27	2.59	3.79	3.7	*		-	-
PLACE1005232	37.45	31.95	45.29	20.08	17.68	11.81	24.42	25.92	23.95	**	*	-	-
PLACE1005238	7.93	6.66	12.45	5.29	3.68	4.36	3.81	3.89	4.88			-	-
PLACE1005243	16.13	27.58	27.13	19.28	15.8	19.26	6.8	12.48	9.67	*		-	-
PLACE1005260	5.37	6.75	7.77	5.96	4.93	3.55	3.78	5.49	3.24			-	-
PLACE1005261	61.12	61.95	56.67	47.53	38.14	42.82	20.88	23.52	21.14	**	**	-	-
PLACE1005266	7.03	6.91	9.71	5.82	8.14	5.96	4.18	3.59	5.53	*		-	-
PLACE1005271	8.47	8.61	6.86	4.77	7.74	3.27	3.85	3.67	3.35	**		-	-
PLACE1005277	20.82	15.1	22.31	9.09	13.8	9.19	14.68	15.84	18.36	*	*	-	-
PLACE1005287	15.1	8.12	14.31	5.25	6.44	6.19	5.01	4.78	3.02	*	*	-	-
PLACE1005289	19.04	8.09	8.79	4.5	5.75	7.77	4.03	4.98	7.58			-	-
PLACE1005305	8.97	7.38	9.77	4.47	3.45	4.18	3.14	3.99	3.81	**	**	-	-
PLACE1005307	12.01	9.78	33.93	6.5	8.01	6.92	5.78	6.91	8.37			-	-
PLACE1005308	6.72	9.28	13.18	4.37	5.39	4.05	5.69	4.54	7.02			-	-
PLACE1005313	8.46	6.04	10.79	5.16	6.95	3.74	4.51	3	4	*		-	-
PLACE1005320	9.29	12.54	14.31	7.32	9.29	5.14	4.92	4.65	5.56	**		-	-
PLACE1005327	31	25.4	35.29	16.09	17.4	11.88	9.82	14.36	14.83	**	**	-	-
PLACE1005331	13.95	13.36	13.69	9.09	7.58	7.27	7.77	10.5	7.38	**	**	-	-
PLACE1005335	11.38	7.84	13.28	5.94	7.87	3.68	8.16	25.07	9.78			-	-
PLACE1005336	29.59	23.98	23.59	14.99	18	12.79	23.63	28.14	27.34	*		-	-
PLACE1005351	11.98	8.58	19.03	4.6	4.51	3.68	3.27	8.24	4.61	*		-	-
PLACE1005366	73.8	70.81	70.61	45.9	65.72	42.52	55.33	53.05	50.72	*	**	-	-
PLACE1005373	9.8	5.9	15.61	3.87	3.73	2.69	2.9	3.58	2.99	*	*	-	-
PLACE1005374	31.25	37.69	80.27	12.29	20.74	12.12	33.17	26.39	29.29	**		-	-
PLACE1005383	9.15	7.81	8.27	5.38	5.79	4.82	3.77	5.77	7.02	*		-	-
PLACE1005388	14.78	7.34	14.13	9.31	7.11	8.02	8.28	5.49	5.89			-	-
PLACE1005409	11.88	10.31	14.09	4.95	5.84	5.2	5.09	4.29	5.35	**	**	-	-
PLACE1005410	13.35	8.74	14.55	5.14	6.18	4.53	5.85	5.79	5.13	**	*	-	-
PLACE1005426	10.65	4.8	11.31	4.23	3.51	3	2.46	6.40	1.78			-	-
PLACE1005431	8.27	7.3	10.2	3.04	3.82	2.32	3.59	3.79	5.31	**	*	-	-
PLACE1005453	7.65	9.23	18.95	3.09	5.38	3.83	2.3	4.42	3.43			-	-
PLACE1005467	10.75	11.01	23.28	4.33	7.93	7.48	6.29	7.38	5.29			-	-
PLACE1005471	7.1	9.43	5.6	3.8	6.04	3.15	3.31	6.18	12.95			-	-
PLACE1005476	29.52	27.39	28.6	15.67	12.74	15.87	14.04	14.17	18.82	**	**	-	-
PLACE1005477	8.37	7.91	11.74	3.68	6.31	4.63	5	3.74	6.24	*	*	-	-
PLACE1005480	7.1	8	8.36	5.81	4.81	4.45	4.92	2.94	2.27	**	**	-	-
PLACE1005481	10.21	6.44	10.18	3.68	4.76	4.41	5.76	4.83	3.43	*	*	-	-
PLACE1005494	9.19	9.94	10.69	3.34	4.05	3.47	4.87	3.69	4.02	**	**	-	-

126 / 157

2691

2692

PLACE1005495	11.05	11.48	9.86	5.01	5.38	4.28	3.3	3.97	3.58	**	**	-	-
PLACE1005497	9.89	8.26	8.02	3.87	6.07	3.47	3.99	3.58	2.73	**	**	-	-
PLACE1005499	23.48	21.47	29.68	8.85	19.6	10.05	16	19.54	21.59	*	*	-	-
PLACE1005502	9.18	6.48	6.37	3.45	3.87	4.29	4.21	4.81	4.94	*	*	-	-
PLACE1005513	32.31	27.28	25.19	10.65	9.45	7.45	5.15	5.68	9.33	**	**	-	-
PLACE1005515	11.7	9.47	10.48	3.66	3.88	4.7	6	4.7	5.35	**	**	-	-
PLACE1005519	8.43	6.62	7.72	4.49	2.77	2.61	3.1	2.38	5.05	**	*	-	-
PLACE1005528	8.8	10.4	11.16	2.78	4.21	2.23	3.21	3.51	3.18	**	**	-	-
PLACE1005528	26.81	22.55	23.67	12.32	10.17	8.61	19.08	11.09	17.43	**	*	-	-
PLACE1005530	20.34	15.73	21.3	12.14	14.27	8.82	8.49	8.49	4.49	*	*	-	-
PLACE1005530	18.12	14.92	18.89	3.17	4.7	3.13	5.93	4.22	4.18	**	**	-	-
PLACE1005539	42.25	35.31	35.69	15.02	20.49	15.33	32.11	27.06	37.95	**	*	-	-
PLACE1005543	35.88	20.89	14.02	11.25	7.4	11.67	9.86	8.84	11.47	*	*	-	-
PLACE1005544	4.88	5.73	4.38	2.81	2.15	3.58	3.84	2.43	3.09	*	*	-	-
PLACE1005550	26.58	26.83	19.78	16.15	16.36	18.32	11.17	10.41	11.67	*	**	-	-
PLACE1005554	23.49	15.23	16.27	7.02	8.22	4.71	11.04	7.45	10.08	*	*	-	-
PLACE1005557	10.41	9.97	10.58	3.15	3.37	2.65	4.57	1.9	2.76	**	**	-	-
PLACE1005563	10.87	11.48	9.72	4.64	6.41	6.88	7.08	5.98	6.9	**	**	-	-
PLACE1005569	27.85	17.24	18.24	8.47	8.97	6.39	8.6	5.75	9.27	*	*	-	-
PLACE1005674	29.21	21.81	10.09	12.87	15.06	14.88	14.12	11.93	13.7	*	*	-	-
PLACE1005684	60.18	86.17	89.3	73.19	89.15	68.42	53.92	40.06	44.98	*	*	-	-
PLACE1005690	14.81	9.97	11.58	8.51	9.72	8.47	7.14	6.35	7.74	*	*	-	-
PLACE1005695	23.18	18.26	21.18	12.01	10.55	9.94	18.07	11.17	15.81	**	*	-	-
PLACE1005801	11.31	7.51	15.39	5.61	6.28	8.37	7.21	9.02	6.01	*	*	-	-
PLACE1005803	18.98	21.86	14.52	8.39	9.87	6.77	12.21	11.52	14.3	*	*	-	-
PLACE1005804	8.92	8.05	5.88	5	4.08	4.87	3.54	2.41	2.72	*	**	-	-
PLACE1005811	33.68	22.2	29.76	11.8	16.07	9.26	20.54	21.42	26.07	*	*	-	-
PLACE1005822	30.69	27.64	23.31	11.83	18.8	12.91	22.33	27.88	24.08	*	*	-	-
PLACE1005823	9.95	5.8	5.03	2.69	3.21	2.48	4.37	5.18	4.84	*	*	-	-
PLACE1005830	8.17	9.66	9.09	4.43	7.2	3.43	4.75	3.89	4.7	*	**	-	-
PLACE1005839	60.6	68.64	49.38	28.59	34.73	29.78	61.43	56.47	58.03	**	**	-	-
PLACE1005846	17.8	13.84	15.64	9.47	9.48	7.41	6.37	4.37	5.79	**	**	-	-
PLACE1005847	13.8	11.09	12.51	5.68	5.78	5.71	6.46	4.42	5.24	**	**	-	-
PLACE1005849	28.95	31.24	20.78	11.15	9.99	10.48	23.48	13.79	10.43	**	*	-	-
PLACE1005853	18.03	13.03	11.9	6.52	4.03	5.5	8.65	5.02	7.5	**	*	-	-
PLACE1005855	15.9	11.93	10.98	7.3	8.49	5.69	13.9	11.98	12.97	*	*	-	-
PLACE1005859	10.6	8.34	6.31	5.3	4.01	3.82	4.61	3.03	4.74	*	*	-	-
PLACE1005860	12.82	10.45	9.32	4.78	5.95	3.56	5.23	2.28	5.04	**	**	-	-
PLACE1005864	23.43	27.28	25.73	6.88	20.98	19.65	21.23	13.1	15.7	*	*	-	-
PLACE1005868	12.43	11.64	13.67	7.79	18.52	6.89	11.1	12.9	13.47	*	*	-	-
PLACE1005869	14.08	13.18	10.88	6.38	8.32	3.2	8.01	5.64	11	*	*	-	-
PLACE1005882	22.57	20.73	18.81	7.71	18.55	7.1	12.18	6.07	11.14	**	**	-	-
PLACE1005898	7.18	6.68	8.02	7.04	5.88	4.02	4.11	5.29	4.18	**	*	-	-
PLACE1005708	17.69	17.02	22.85	11.44	15.51	9.8	12.69	8.54	8.25	*	*	-	-
PLACE1005725	9.73	11.27	13.87	7.08	8.31	6.27	4.4	6.48	6.4	*	*	-	-
PLACE1005727	10.04	8.34	11.08	6.79	18.78	10.55	4.59	6.83	6.58	*	*	-	-
PLACE1005750	13.44	12.88	11.77	6.36	7.19	8.47	6.04	7.92	8.47	**	**	-	-
PLACE1005756	5.83	5.5	7.74	3.15	14.38	1.73	4.63	3.09	3.33	*	*	-	-
PLACE1005759	7.01	13.06	9.75	7.43	7.8	5.14	2.98	4.18	3.47	*	*	-	-
PLACE1005745	8.04	6.14	6.43	4.28	4.82	5.89	4.13	2.83	4.39	*	*	-	-
PLACE1005752	128.38	102.07	139.38	89.6	72.88	89.83	103.85	94.56	103.55	*	*	-	-
PLACE1005755	13.48	9.22	14.54	4.95	8.97	3.84	5.58	5.85	3.87	*	*	-	-
PLACE1005756	10.54	15.55	18.58	10.11	10.59	10.1	6.71	10.57	12.53	*	*	-	-
PLACE1005760	5.35	5.71	6.64	9.33	6.58	7.47	2.74	8.03	2.74	*	*	-	-
PLACE1005763	18.11	17.1	15.35	9.45	11.68	9.73	7.29	9.28	9.82	**	**	-	-
PLACE1005768	19.31	21.05	13.08	10.64	11.47	13.73	6.13	9.41	9.52	*	*	-	-
PLACE1005771	24.23	23.82	24.14	14.88	21.27	9.53	16.05	19.49	20.2	*	*	-	-
PLACE1005783	6.96	4.13	7.26	3.95	3.94	3.49	3.16	3.04	5.15	*	*	-	-
PLACE1005790	43.95	40.56	57.56	28.57	25.88	15.98	30.85	28.7	32.18	*	*	-	-
PLACE1005802	18.73	15.75	12.71	10.85	14.22	15.85	21.48	13.68	14.98	*	*	-	-
PLACE1005803	14.16	12.88	18.69	6.94	7.08	7.31	4.95	5.48	3.9	**	**	-	-
PLACE1005804	16.14	18.08	21.87	10.88	8.62	11.88	4.49	5.03	5.63	*	**	-	-
PLACE1005813	7.59	9.87	17.86	7.9	9.25	7.77	3.88	4.99	5.43	*	*	-	-
PLACE1005815	17.11	18.5	18.61	7.94	18	6.8	9.57	8.11	8.77	**	*	-	-

127 / 157

【0951】

【表480】

2694

PLACE1005820	21.32	17.86	17.66	18.38	28.15	11	8.41	12.77	17.42				
PLACE1005833	19.04	15.23	17.55	13.68	13.86	12.18	11.62	13.97	12.82				
PLACE1005834	21.23	40.41	42.81	19.53	15.14	12.45	22.97	22.37	27.18	00	0	-	-
PLACE1005835	8.38	4.15	8.95	3.69	3.05	3.26	2.72	13.93	4.26				
PLACE1005836	6.34	4.47	7.11	2.95	2.55	2.1	1.11	4.41	2.67	*	*	-	-
PLACE1005845	7.96	5.73	10.94	3.26	4.7	1.75	3.2	4.47	3.82	*	*	-	-
PLACE1005850	7.12	8.61	11.98	4.69	4.51	3.39	2.96	4.97	6.83	*	*	-	-
PLACE1005851	49.92	45.48	36.8	26.76	36.45	21.25	9.27	9.06	15.36		00	0	-
PLACE1005856	9.13	9.45	9.88	4.42	6.52	4.26	5.86	9.06	7.88	*	*	-	-
PLACE1005875	19.11	12.98	14.88	10.98	11.98	11.34	7.25	9.07	9.2	*	*	-	-
PLACE1005876	6.84	5.72	8.28	3.52	3.77	3.26	2.47	4.32	3	*	*	-	-
PLACE1005878	6.18	5.12	8.15	3.65	4.35	1.9	2.1	7.39	4.67				
PLACE1005880	10.13	10.1	15.44	15.29	7.51	5.59	11.17	12.56	9.82				
PLACE1005884	15.13	11.77	15.14	4.99	6.58	3.8	5.03	3.76	7.31	**	00	-	-
PLACE1005880	12.42	12.54	15.15	4.55	6.88	4.95	4.84	4.65	10.58	00	0	-	-
PLACE1005888	24.17	43.11	28.69	12.33	15.21	13.1	10.14	6.45	7.84	*	*	-	-
PLACE1005913	11.08	12.76	18.39	6.03	9.77	5.43	8.89	7.87	6.55				
PLACE1005921	38.7	26.83	45.11	14.95	20.53	13.83	14.46	15.3	19.69	*	*	-	-
PLACE1005923	35.55	28.8	35.14	8.99	9.9	10.33	7.55	9.92	9.16	**	00	-	-
PLACE1005925	9.28	6.52	9.89	4.22	6.78	3.82	4.38	4.97	3.85	*	*	-	-
PLACE1005927	8.32	7.66	16.91	2.68	3.86	2.87	2.78	4.08	4.5				
PLACE1005932	17.99	14.04	15.58	5.88	10.87	7.85	11.54	8.65	8.07	*	*	-	-
PLACE1005934	7.66	7.62	8.85	2.45	3.18	2.78	2.65	3.47	4.68	**	00	-	-
PLACE1005935	8.91	7.48	10.96	2.6	4	2.78	2.84	2.57	2.62	**	00	-	-
PLACE1005939	14.59	14.49	13.88	5.4	8.87	6.89	7.24	6.14	6.85	**	00	-	-
PLACE1005951	164.11	138.22	202.24	63.95	124.2	100.53	168.66	124.78	76.11	*	*	-	-
PLACE1005953	4.12	2.98	5.89	2.81	3.85	1.89	1.88	2.73	1.8				
PLACE1005955	147.21	90.51	183.74	157.35	162.68	147.78	83.65	73.22	67.87				
PLACE1005966	21.68	17.57	18.88	12.22	12.03	11.48	18.97	16.96	19.33	**	00	-	-
PLACE1005988	7.85	6.88	6.59	3.18	3.12	3.27	2.21	3.25	2.88	**	00	-	-
PLACE1005975	6.58	6.62	8.7	2.58	2.3	2.44	2.19	2.22	2.24	**	00	-	-
PLACE1005980	13.31	13.63	10.46	5.9	8.18	4.45	5.05	6.3	5.08	**	00	-	-
PLACE1005987	11.27	8.84	8.81	3.34	7.88	4.32	3.83	4.17	2.65	**	00	-	-
PLACE1006002	7.2	9.3	8.82	4.13	8.01	2.96	3.03	3.74	3.08	**	00	-	-
PLACE1006003	19.13	15.27	13.88	6.84	9.41	7.04	14.37	14.53	15.21	*	*	-	-
PLACE1006011	14.53	12.13	13.55	5.17	6.37	5.39	12.92	13.32	12.8	**	00	-	-
PLACE1006017	17.88	15.33	13.64	7.8	8.85	7.28	13.44	3.4	5.68	**	00	-	-
PLACE1006037	19.64	15.54	16.94	5.06	9.69	7.83	15.52	15.45	11.65	**	00	-	-
PLACE1006040	21.73	18.19	20.02	8.22	10.07	7.19	9.12	8.1	8.15	**	00	-	-
PLACE1006083	18.04	22.51	29.29	15.12	25.84	16.72	12.74	9.35	9.64	*	*	-	-
PLACE1006071	24.45	22.47	19.69	9.7	14.01	7.6	15.44	10.65	16.88	**	0	-	-
PLACE1006073	38.01	25.19	30.05	16.58	18.78	14.04	20.88	12.98	14.52	*	*	-	-
PLACE1006074	21.68	18.68	21.89	13.65	9.89	8.37	10.34	8.49	11.15	**	00	-	-
PLACE1006076	6.98	23.32	28.44	8.28	24.88	6.88	8.27	11.5	12.4				
PLACE1005979	5.38	7.69	6.82	3.43	3.4	2.8	2.82	1.69	5.77	*	*	-	-
PLACE1006082	17.29	20.65	15.5	11.28	11.48	8.91	6.51	3.34	7.24	*	*	-	-
PLACE1006116	6.88	4.88	6.43	2.05	2.48	1.97	3.15	0.18	1.1	**	0	-	-
PLACE1006119	9.01	6.23	8.78	2.16	3.24	2.75	3.83	1.45	4.03	**	0	-	-
PLACE1005129	11.8	19.99	12.49	6.74	8.91	5.37	6.93	6.8	9.62	*	*	-	-
PLACE1004139	11.03	9.2	11	5.5	4.05	3.64	5.37	5.2	8.18	**	0	-	-
PLACE1006143	34.11	32.02	28.21	13.95	17.22	16.38	5.72	10.56	8.1	*	*	-	-
PLACE1006157	22.08	18.11	21.21	6.21	8	10.68	9.26	8.55	9.67	**	00	-	-
PLACE1006158	22.74	19.17	16.53	5.42	7.39	5.18	11.09	3.24	9.93	**	0	-	-
PLACE1006164	10.4	12.06	12.25	3.81	6.89	2.98	4.98	2.08	3.84	**	00	-	-
PLACE1006167	150.43	130.54	128	185.79	82.45	55.64	118.18	93.17	93.5	*	*	-	-
PLACE1006170	12.04	9.61	11.88	3.48	3.73	3.68	2.75	2.82	3.04	**	00	-	-
PLACE1006181	14.32	18.19	10.38	5.74	6.23	4.42	6.29	2.91	4.78	**	0	-	-
PLACE1006187	29.88	21.9	18.48	7.26	12.29	5.92	18.37	17.04	21.08	**	00	-	-
PLACE1006195	27.43	19.87	20.2	12.18	6.92	6.66	10.17	6.16	6.59	*	*	-	-
PLACE1006196	14.88	10.51	8.01	6.22	7.31	2.89	4.66	4.93	4.37	*	*	-	-
PLACE1006197	25.54	27.48	23.65	11.8	18.78	16.24	17.7	18.29	21.21	*	*	-	-
PLACE1006198	12.16	16.37	13.48	4.05	6.5	3.12	5.11	3.17	4.8	**	00	-	-
PLACE1006205	8.38	7.44	9.1	4.62	5.35	3.49	12.53	5	7.63	**	00	-	-
PLACE1006208	10.95	10.22	8.53	5.04	9.25	5.28	4.47	4.13	4.33	**	00	-	-

2695

2696

PLACE1006211	18.16	19.48	15.85	7.8	6.58	8.16	10.56	8.16	8.88	**	**	-	-
PLACE1006219	28.22	24.03	22.82	13.99	7.2	9.8	23.91	20.19	22.03	**	**	-	-
PLACE1006221	33.12	26.98	22.85	14.89	11.88	8.81	23.85	22.74	28.77	**	**	-	-
PLACE1006225	9.38	7.86	8.1	4.18	4.91	3.81	6.34	5.77	7.87	*	*	-	-
PLACE1006230	23.05	26.18	25.58	13.5	14.02	10.31	12.5	8.37	11.47	**	**	-	-
PLACE1006239	7.77	5.03	6	2.21	3.14	2.06	2.98	3.13	3.81	*	*	-	-
PLACE1006245	11.97	10.97	10.73	4.72	5.26	4.25	5.6	4.82	4.52	**	**	-	-
PLACE1006248	14.07	10.52	9.12	4.17	12.12	4.44	10.44	6.15	11.84	*	*	-	-
PLACE1006248	10.3	9.17	8.58	6.54	7.05	2.85	7.87	4.65	5.48	*	*	-	-
PLACE1006252	19.03	26.43	25.68	10.77	26.8	17.9	14.07	10.91	15.08	*	*	-	-
PLACE1006259	6.35	4.8	5.62	2.78	2.48	2.42	4.23	3.87	2.13	**	*	-	-
PLACE1006275	7.84	6.83	8.11	7.79	9.48	8.48	5.82	11.38	4.94	*	*	-	-
PLACE1006277	18.18	15.2	14.16	12.95	19.85	15.77	6.89	13.62	7.81	*	*	-	-
PLACE1006288	23.16	25	24.73	14.28	21	15.27	15.12	10.66	20.01	*	*	-	-
PLACE1006290	3.79	4.35	4.22	1.07	3.07	1.54	0.78	2.83	1.89	*	*	-	-
PLACE1006298	31.99	33.57	40.35	21.82	22.17	18.83	30.18	28.42	31.31	**	**	-	-
PLACE1006311	24.08	25.54	26.15	17.19	20.35	18.87	6.49	6.41	12.35	**	**	-	-
PLACE1006318	23.92	22.31	22.74	9.17	11.04	10.73	16.57	17.39	15.35	**	**	-	-
PLACE1006325	10.97	8.18	13.93	5.64	6.43	6.48	6.34	4.38	3.68	*	*	-	-
PLACE1006331	322.49	249.29	312.14	284.34	344.95	317.78	281.41	376.24	187.88	*	*	-	-
PLACE1006335	6.01	6.97	9.42	6.43	10.77	6.27	4.79	4.63	3.75	*	*	-	-
PLACE1006357	23.6	23.65	28.41	15.45	19.82	11.71	10.11	13.97	13.07	*	**	-	-
PLACE1006360	39.03	55.89	60.83	32.58	29.33	38.42	18.81	22.42	24.9	*	*	-	-
PLACE1006364	4.49	3.79	4.52	5.2	6.32	3.04	3.31	4.72	3.85	*	*	-	-
PLACE1006365	8.87	13.33	7.94	4.35	7.31	4.11	4.93	6.19	6.08	*	*	-	-
PLACE1006368	7.6	5	7.22	4.35	3.51	2.72	3.12	4.28	3.68	*	*	-	-
PLACE1006371	12.84	8.89	13.13	6.85	5.51	3.89	5.93	6.19	7.38	*	*	-	-
PLACE1006373	31.88	31.57	27.89	11.44	15.9	12.05	24.23	17.9	24.85	**	*	-	-
PLACE1006382	15.81	18.11	17.64	20.83	24.97	21.9	28.68	49.67	48.66	*	+	+	-
PLACE1006385	23.74	24.67	19.54	18.57	28.47	23.59	6.72	12.92	8.24	**	*	-	-
PLACE1006391	7.09	8.24	10.32	5.78	7.9	5.35	7.26	7.31	6.16	*	*	-	-
PLACE1006412	14.35	15.98	15.55	14.85	16.87	8.5	8.42	8.73	11.54	**	*	-	-
PLACE1006414	30.84	83.89	32.69	29.17	49.53	18.72	7.89	13.37	10.44	*	*	-	-
PLACE1006419	23.86	16.38	19.21	8.34	10.82	5.22	6.84	8.36	7.78	*	**	-	-
PLACE1006438	16.88	12.76	13.77	5.93	7.83	5.98	6.47	9.15	5.41	**	*	-	-
PLACE1006443	7.62	8.28	7.5	4.32	3.07	4.42	2.72	3.77	2.68	**	**	-	-
PLACE1006445	53.01	52.47	65.34	48.09	51.7	50.7	24.1	28.82	30.98	**	*	-	-
PLACE1006447	18.46	17.73	17.85	9.84	12.81	9.03	5.67	7.16	5.82	**	**	-	-
PLACE1006466	5.01	5.78	10.4	3.5	5.28	3.03	3.52	2.47	3.82	*	*	-	-
PLACE1006469	9.97	6.93	12.7	4.45	5.94	2.89	6.14	6.08	3.89	*	*	-	-
PLACE1006470	14.91	13.1	13.03	5.44	14.24	9.05	12.85	15.12	16.52	*	*	-	-
PLACE1006472	32.56	24.9	52.53	14.88	22.47	15.58	27.13	60.66	18.13	*	*	-	-
PLACE1006476	75.24	88.04	85	49.83	68.66	43.58	73.04	95.42	77.62	*	*	-	-
PLACE1006482	19.52	14.87	22.77	7.46	8.75	5.8	9.29	24.88	15.09	**	*	-	-
PLACE1006488	13.42	9.51	14.11	5.93	5.58	7.22	5.43	7.77	6	*	*	-	-
PLACE1006492	53.41	60.91	85.09	29.78	38.14	28.4	41.31	52.42	43.57	**	*	-	-
PLACE1006505	8.83	8.04	10.86	3.97	6.16	3.99	2.77	5.01	4.34	*	**	-	-
PLACE1006515	6.57	6.17	10.41	2.89	4.8	3.13	3.49	3.57	2.51	*	*	-	-
PLACE1006518	7.77	7.4	8.18	2.65	6.76	3.8	4.8	4.81	1.69	*	*	-	-
PLACE1006520	85.74	60.72	84.5	67.78	73.09	68.07	40.48	39.38	37.31	**	*	-	-
PLACE1006521	5.4	3.13	7.19	3.67	3.58	2.13	2.56	2.69	1.53	*	*	-	-
PLACE1006529	21.37	15.93	24.45	11.65	12.23	11.32	13.1	15.15	18.91	*	*	-	-
PLACE1006531	8.88	4.63	8.48	3.15	2.87	3.1	2.57	4.89	2.69	*	*	-	-
PLACE1006534	11.24	12.38	19.26	6.83	8.12	7.2	8.08	5.98	8.82	*	*	-	-
PLACE1006540	17.3	13.48	19.89	8	8.01	6.33	5.04	4.77	5.56	**	**	-	-
PLACE1006549	67.43	78.29	59.13	33.67	55.25	39.07	49.85	38.47	58.34	*	*	-	-
PLACE1006550	27.31	23.34	17.69	10.6	24.88	11.76	7.55	11.68	6.46	*	*	-	-
PLACE1006552	41.43	35.42	38.21	18.37	17.9	12.42	28.67	34.48	34.92	**	*	-	-
PLACE1006557	68.91	37.28	57.05	36.71	38.26	34.22	31.47	33.21	28.65	*	*	-	-
PLACE1006563	12.53	9.16	14.73	7.21	10.95	7.06	5.43	4.33	7.28	*	*	-	-
PLACE1006579	8.16	5.5	6.78	3.58	3.74	3.1	2.74	1.94	3.28	*	**	-	-
PLACE1006594	8.47	4.75	7.38	2.29	3.19	1.42	2.51	1.45	1.95	*	*	-	-
PLACE1006598	14.33	13.22	15.98	9.86	13.18	9.8	7.97	4.5	8.78	**	*	-	-
PLACE1006607	18.95	13.98	15.61	10.13	11.38	9.38	8.88	8.34	9.08	*	**	-	-

129 / 157

2697

2698

PLACE1006810	9.23	9.13	6.03	3.58	3.78	2.24	3.21	1.79	1.8	**	**	-	-
PLACE1006815	5.13	4.58	7.74	3.64	4.24	3.19	3.42	3.13	4.2			-	-
PLACE1006817	10.84	9.51	10.84	5.44	6.44	4.54	8.05	7.65	8.94	**	*	-	-
PLACE1006818	16.8	17.21	17.68	12.4	9.99	12.65	9.98	9.85	12.85	**	**	-	-
PLACE1006826	9.9	9.38	11.8	4.89	3.64	3.15	3.8	3.9	3.22	**	**	-	-
PLACE1006829	97.87	77.5	62.9	75.16	69.13	71.45	50.39	48.85	45	*		-	-
PLACE1006837	20.62	30.53	27.31	15.03	24.3	12.77	12.77	11.41	9.69	**		-	-
PLACE1006840	148.09	116.55	121.82	98.73	144.79	100.85	85.85	81.12	81.72	**		-	-
PLACE1006844	10.39	13.78	11.01	6.08	7.82	8.85	6.47	3.27	4.59	*		-	-
PLACE1006857	9.47	9.7	8.11	3.37	3.88	4.59	4.46	4.41	6.15	**	**	-	-
PLACE1006873	128.37	101.12	117.47	60.38	47.28	69.57	78.35	78.65	91.83	**	*	-	-
PLACE1006878	13.14	7.54	6.57	3.78	6.24	6.14	3.95	3.22	3.7			-	-
PLACE1006882	5.12	3.88	4.9	2.21	2.14	1.84	2.5	1.43	2.09	**	**	-	-
PLACE1006884	8.48	6.37	7.5	1.9	2.47	2.03	3.67	2.07	2.03	**	**	-	-
PLACE1006888	24.85	21.3	17.68	6.94	8.35	11.27	7.31	3.9	5.72	**	**	-	-
PLACE1006904	10.76	10.1	9.67	7.04	6.45	4.58	4.88	2.24	3.52	**	**	-	-
PLACE1006908	6.82	6.82	4.54	1.43	4.58	2.49	4.44	1.85	2.6	*		-	-
PLACE1006911	7.97	5.29	8.07	5.12	5.44	6.82	5.18	4.48	3.42			-	-
PLACE1006914	19.78	16.32	14.11	6.39	7.42	10.52	4.48	5.02	4.69	*	**	-	-
PLACE1006916	21.34	19.47	16.01	5.39	7.72	6.97	12.01	9.74	13.28	**	*	-	-
PLACE1006921	8.78	6.59	9.88	4.18	4.28	2.45	3.4	4.54	3.95	*	*	-	-
PLACE1006954	12.83	10.28	11.39	6.83	6.2	6.24	8.13	4.13	6.9	**	*	-	-
PLACE1006960	17.66	24.39	16.04	22.5	23.48	11.75	21.46	9.03	14.39	*	*	-	-
PLACE1006979	14.53	14.99	8.23	6.82	4.21	3.38	5.07	3.29	2.35	*	*	-	-
PLACE1006982	13.34	14.85	8.6	8.18	13.03	7.71	10.87	6.95	6.98			-	-
PLACE1006983	28.1	20.43	20.94	18.8	23.33	15.04	25.48	18.52	26.77			-	-
PLACE1006986	16.5	11.01	11.97	10.81	6.14	7.77	7.76	8.87	8.16	*		-	-
PLACE1006992	43	29.8	28.89	12.14	32.07	8.88	14.31	11.42	11.6	*		-	-
PLACE1006995	6.48	5.54	6.42	3.27	4.53	2.37	5.98	1.9	3.55	*		-	-
PLACE1006800	14.69	12.28	14.82	7.42	7.88	5.38	11.02	8.83	12.6	**		-	-
PLACE1006805	297.4	379.55	330.72	261.58	242.8	138.41	164.1	78.29	121.42	*	**	-	-
PLACE1006809	17.57	15.02	14.01	6.89	5.54	5.55	12.59	5.87	9	**	*	-	-
PLACE1006815	38.78	37.15	24.87	22.04	27.53	19.08	12.45	10.82	14.81	**	**	-	-
PLACE1006819	8.55	6.96	7.82	3.54	3.27	1.58	3.56	2.07	3.34	**	**	-	-
PLACE1006820	11.34	11.23	10.69	7.28	7.6	3.88	8.72	6.8	7.48	*	**	-	-
PLACE1006825	11.58	10.21	8.37	4.28	6.11	3.55	4	5.42	6.14	*	*	-	-
PLACE1006829	8.68	7.12	6.95	4.95	4.5	3.35	3.73	4.63	3.79	*	**	-	-
PLACE1006853	7.78	9.04	10.08	4.43	6.8	3.94	5.65	3.81	3.65	**	**	-	-
PLACE1006860	13.3	14.13	11.82	7.29	7.57	5.64	7.39	5.43	7.17	**	**	-	-
PLACE1006867	20.52	19.85	17.03	7.58	12.2	4.52	8.89	4.5	9.57	*	**	-	-
PLACE1006875	6.86	5.29	3.88	1.69	3.65	1.95	4.84	3.37	3.97	*		-	-
PLACE1006878	5.96	7.5	9.4	3.45	4.02	4.56	4.29	5.32	7.08	*		-	-
PLACE1006883	17.58	24.05	26.14	11.43	14.88	9.68	8.41	13.21	13.16	*	*	-	-
PLACE1006898	34.11	35.72	38.43	15.89	26.64	17.27	29.81	38.71	32.24	*		-	-
PLACE1006901	35.59	25.34	29.52	17.57	24.59	23.14	7.08	14.35	10.28	**		-	-
PLACE1006904	283.91	237.88	285.1	210.53	264.05	203.01	176.82	124.88	215.9			-	-
PLACE1006917	8.87	8.87	10.42	4.77	9.67	4.27	2.95	3.43	5.44	**		-	-
PLACE1006932	13.9	19.05	15.18	17.43	14.78	12.47	12.23	11.92	6.34			-	-
PLACE1006936	6.15	5.58	5.5	4.25	5.8	3.94	4.86	3.59	4.94			-	-
PLACE1006956	73	65.43	79.54	37.1	42.04	37.7	45.33	35.06	47.75	**	**	-	-
PLACE1006958	14.71	17.1	17.15	7.75	10.57	5.67	10.62	14.42	12.92	**		-	-
PLACE1006959	8.43	6.88	7.08	3.86	5.4	3.09	3.53	7.7	6.81	*		-	-
PLACE1006981	11.78	13.88	26.99	6.58	7.47	6.43	7.89	10.93	8.94			-	-
PLACE1006982	8.5	7.71	10.43	6.5	8.54	6.61	4.94	6	6.79	*		-	-
PLACE1006986	30.69	83.35	44.81	30.7	31	25.81	17.51	20.26	25.7			-	-
PLACE1006978	9.85	10.58	9.97	5.88	6.1	5.13	5.32	5.52	6.47	**	**	-	-
PLACE1006989	23.14	43.35	32.93	27.05	47.4	17.28	23.49	21.59	18.62			-	-
PLACE1007001	14.41	12.68	17.15	10.97	5.54	6.74	6.74	7.87	8.22	*	**	-	-
PLACE1007014	9.24	9.35	10.83	6.4	8.04	5.94	4.98	6.15	5.52	**	**	-	-
PLACE1007021	66.53	38.34	63.48	20.03	40.71	38.97	20.72	29.77	27.83	*		-	-
PLACE1007028	85.54	56.8	77.58	57.55	60.7	57.93	57.5	56.64	67.29			-	-
PLACE1007028	8.5	10.78	17.58	7.89	7.82	6.28	3.31	5.38	3.91	*		-	-
PLACE1007038	7.88	7.89	13.88	4.19	6.33	3.79	2.05	3.52	4.03	*		-	-
PLACE1007040	6.13	9.51	15.11	4.74	7.09	4.87	8.84	8.35	6.98	*		-	-

130 / 157

2699

2700

PLACE1007045	16.78	21.85	22.19	13.09	30.39	11.83	9.78	12.74	10.75	*	*	-
PLACE1007046	19.3	14.44	24.23	9.41	10.57	11.83	9.91	12.06	15.93	*	*	-
PLACE1007053	9.31	5.29	12.35	5.07	3.06	4.58	5.36	25.19	5.41	*	*	-
PLACE1007068	8.87	4.77	10.34	5.02	4.28	3.54	2.63	5.02	2.74	*	*	-
PLACE1007070	18.55	13.15	15.29	11.03	10.78	10.43	6.96	7.21	5.63	*	**	-
PLACE1007076	17.03	17.73	24.8	24.48	28.32	20.68	16.51	16.83	13.48	*	*	-
PLACE1007077	7.35	10.92	12.28	4.54	5.11	3.6	8.43	5.8	5.66	*	*	-
PLACE1007081	41.38	35.79	51.18	37.04	46.87	29.31	28.18	20.46	26.31	*	*	-
PLACE1007082	14.74	14.27	12.93	7.51	10.12	6.83	13.51	10.86	10.42	*	*	-
PLACE1007092	9.52	6.79	8.77	4.99	5.23	4.04	4.42	5.84	7.46	*	*	-
PLACE1007095	8.7	7.04	11.33	3.9	6.35	2.61	4.31	10.82	7.33	*	*	-
PLACE1007097	5.03	2.47	6.96	1.99	2.48	1.11	1.28	4.01	1.45	*	*	-
PLACE1007099	14.43	9.17	8.68	4.67	6.07	4.14	4.7	5.17	7.04	*	*	-
PLACE1007105	9.93	9.89	19.72	4.22	5.48	3.86	5.83	6.27	9.12	*	*	-
PLACE1007108	7.87	5.2	13.72	3	3.64	1.32	1.96	5.82	6.69	*	*	-
PLACE1007111	24.26	33.98	33.98	8.75	19.55	14.7	10.61	6.46	7.88	*	**	-
PLACE1007112	13.01	15.69	10.39	7.41	17.53	7.88	9.51	4.37	5.03	*	*	-
PLACE1007130	7.28	5.04	7.51	3.97	3.76	4.18	4.15	3.12	4.45	*	*	-
PLACE1007132	11.01	5.03	12.54	6.14	8.23	7.16	4.79	4.84	3.32	*	*	-
PLACE1007140	8.1	8.86	9.35	5.74	3.15	5.4	6.1	3.71	4.08	*	*	-
PLACE1007143	7.85	4.78	7.3	2.78	4.42	3.52	3.05	2.29	1.7	*	*	-
PLACE1007169	21.25	20.42	20.37	9.69	16.47	8.81	13.15	12.99	9.83	*	**	-
PLACE1007178	5.69	6.23	10.18	2.7	4.22	2.51	2.33	2.04	2.65	*	*	-
PLACE1007180	9.72	11.86	11.22	5.23	6.85	7.11	5.69	6.59	16.1	**	*	-
PLACE1007201	11.31	9.93	8.97	4.82	6.82	3.46	5.63	4.72	7.11	*	*	-
PLACE1007202	14.98	8.83	16.66	5.59	5.66	4.16	5.42	6.71	5.73	*	*	-
PLACE1007226	110.77	93.69	99.41	39.5	91.72	72.21	82.39	66.08	68.98	*	*	-
PLACE1007236	53.4	44.43	54.82	32.05	38.08	27.04	24.59	23.68	27.03	**	**	-
PLACE1007239	22.01	21.57	18.94	9.88	9.77	7.42	7.13	6.76	5.82	**	**	-
PLACE1007242	8.39	6.41	6.58	2.97	3.63	2.45	2.76	1.3	1.2	**	**	-
PLACE1007243	51.6	47.43	46.85	38.48	53.48	36.76	30.03	24.3	28.99	**	*	-
PLACE1007247	13.13	15.22	14.84	6.24	18.32	8.42	5.62	3.59	3.12	*	**	-
PLACE1007257	10	9.05	8.75	2.44	7.1	2.96	4.72	2.48	2.64	*	**	-
PLACE1007274	7.6	6.25	7.77	4.39	3.79	3.63	3.01	2.85	2.82	**	**	-
PLACE1007276	4.51	6.38	6.4	3.01	2.72	2.17	2.35	3.14	6.03	**	*	-
PLACE1007282	6.97	4.05	5.61	2.64	2.11	3.58	3.82	3.22	3.39	*	*	-
PLACE1007286	11.57	9.64	10.32	4.04	8.1	5.17	5.17	3.51	3.72	**	**	-
PLACE1007288	16.68	11.68	9.95	4.46	6.87	6.7	3.08	3.72	1.93	*	*	-
PLACE1007301	14.64	13.62	12.24	3.81	5.07	2.87	5.09	2.86	3.16	**	**	-
PLACE1007314	9.78	10.14	10.9	5.59	4.99	2.5	3.88	2.16	3.2	**	**	-
PLACE1007317	8.02	7.14	6.45	3.78	5.35	2.25	2.6	5.13	4.46	*	*	-
PLACE1007329	6.93	6.45	9.13	3.48	3.47	2.66	3.68	2.3	2.77	**	**	-
PLACE1007338	57.3	54.81	50.58	25.82	27.62	24.83	48.58	43.5	58.48	**	*	-
PLACE1007342	41.9	26.92	32.51	26.5	30.16	25.56	24.13	20.5	25.41	*	*	-
PLACE1007345	5.67	4.07	4.44	3.74	2.2	1.81	1.89	2.29	3.75	*	*	-
PLACE1007346	20.5	16.07	16.53	8.68	6.27	3.82	4.58	3.49	4.29	**	**	-
PLACE1007359	41.26	41.58	38.73	21.18	26.14	23.96	27.05	27.12	33.74	**	**	-
PLACE1007367	8.85	6.31	7.86	1.51	3.49	1.31	3.83	1.42	3.11	**	**	-
PLACE1007375	12.55	10.83	9.7	1.39	5.83	1.68	2.54	2.85	2.14	**	**	-
PLACE1007377	6.98	4.2	6.44	4.26	4.62	2.93	3.59	4.11	1.58	*	*	-
PLACE1007386	12.07	10.56	14.15	4.68	4.55	5.21	6.12	7.62	8.27	**	*	-
PLACE1007392	24.21	18.68	19.13	13.92	12.02	7.51	28.83	13.55	12.42	*	*	-
PLACE1007402	16.76	19.32	15.67	8.17	8.78	7.34	11.49	10.03	10.78	**	**	-
PLACE1007409	9.51	9.82	11.04	5.11	5.51	7.23	4.08	2.01	3.77	**	**	-
PLACE1007416	22.54	28.88	22.82	18.84	13.11	6.16	9.22	5.26	7.48	*	**	-
PLACE1007420	14.87	16.22	16.74	8.73	9.19	6.79	10.55	5.47	7.91	**	**	-
PLACE1007431	28.03	25.88	21.18	15.73	16.55	11.3	26.89	15.55	26.27	*	*	-
PLACE1007450	6.82	6.06	4.68	4.08	4.16	1.71	5.65	4.38	4.35	*	*	-
PLACE1007452	14.4	7.71	4.99	5.04	6.12	1.89	4.23	8.8	6.08	*	*	-
PLACE1007454	48.36	48.08	36.43	21.1	38.64	20.65	11.88	16.68	14.49	**	*	-
PLACE1007460	11.52	10.93	12.45	8.16	8.35	4.82	9.8	9.12	12.38	*	*	-
PLACE1007478	12.6	13.38	10.88	4.17	8.18	7.05	7.4	4.59	6.97	*	**	-
PLACE1007484	10.71	8.45	6.68	3.65	4	2.5	4.9	2.97	5.47	*	*	-
PLACE1007488	14.59	14.99	17.16	7.24	4.48	5.55	6.08	3.57	8.77	**	**	-

131 / 157

【0955】

【表484】

2701

2702

PLACE1007507	16.83	14.22	12.4	6.49	10.52	6.63	19.33	11.77	17.21	*	-
PLACE1007511	13.26	11.24	10.12	3.93	3.98	2.6	3.01	2.43	4.23	**	-
PLACE1007513	4.53	2.47	3.69	3.11	1.64	1.97	3.08	2.87	2.04	*	-
PLACE1007524	49.4	41.73	37.68	21.38	27.05	16.04	48.44	42.69	43.18	*	-
PLACE1007525	7.48	5.19	4.73	2.29	4.06	1.12	3.74	3.09	4.26	*	-
PLACE1007537	8.42	7.3	9.61	4.91	6.72	3.83	6.57	5.7	5.93	*	-
PLACE1007544	5.87	4.7	5.81	1.29	3.38	2.37	4.39	1.96	2.74	*	-
PLACE1007547	18.52	13.61	13.32	8.64	13.2	6.36	11.39	8.38	8.52	*	-
PLACE1007557	22.7	28.33	18.93	8.14	13.6	7.63	24.98	15.24	20.74	*	-
PLACE1007560	25.9	29.34	29.57	9.23	13.41	19.28	21.02	29.83	20.17	*	-
PLACE1007566	3.88	2.26	3.15	1.89	4.77	2.57	1.98	2.22	2.43	*	-
PLACE1007580	4.69	3.09	5.83	2.2	6.07	1.42	1.87	5.45	1.54	*	-
PLACE1007583	3.85	2.11	3.18	1.84	4.14	4.12	2.02	2.49	2.5	*	-
PLACE1007591	5.53	6.61	10.65	2.9	4.26	2.95	4.34	4.96	5.37	*	-
PLACE1007598	9.38	9.28	9	2.98	15.1	3.88	2.92	7.29	5.82	*	-
PLACE1007610	3.93	7.09	11.32	4.32	11.34	3.52	4.35	8.81	6.31	*	-
PLACE1007618	4.85	4.13	5.18	3.68	4.39	5.01	3.62	3.16	4.04	*	-
PLACE1007621	20.28	8.89	15.56	6.24	12.9	7.88	10.43	10.08	8.81	*	-
PLACE1007626	35.43	28.05	45.42	25.74	58.46	26.54	38.37	49.51	35.74	*	-
PLACE1007632	8.18	6.81	8.24	5.02	13.18	12.74	5.22	7.37	5.63	*	-
PLACE1007635	17.84	21.72	22.81	15.26	21.82	12.51	7.83	15.31	12.34	*	-
PLACE1007645	29.48	44.32	50.79	29.79	28.69	29.97	18.24	18.23	26.32	*	-
PLACE1007649	5.43	7.88	6.71	4.51	9.94	3.18	4.73	2.18	2.74	*	-
PLACE1007659	23.48	33.01	29.15	18.02	23.66	14.9	25	27.47	23.31	*	-
PLACE1007669	17.27	10.25	11.89	6.18	6.01	5.07	11.02	11.92	11.97	*	-
PLACE1007677	15.96	18.35	15.91	8.53	7.7	12.44	11.15	10.52	8.89	*	-
PLACE1007686	30.37	21.3	31.84	9.3	9.6	8.97	5.43	7.59	4.3	**	-
PLACE1007690	16.29	14.9	13.94	8.07	9.45	10.99	5.55	6.43	7.69	**	-
PLACE1007697	7.22	8.07	6.65	7.42	10.69	12.34	4.08	7.67	5.36	*	-
PLACE1007702	9.68	18.27	14.18	8.71	12.06	12.61	6.79	7.15	7.48	*	-
PLACE1007708	14.91	16.84	14.3	10.72	14.81	6.83	5.13	7.31	8.61	**	-
PLACE1007725	15.22	10.08	11.32	8.01	8.72	6.5	4.92	7.04	8.09	*	-
PLACE1007728	9.62	8.04	8.37	6.72	6.7	9.05	5.49	8.22	6.25	*	-
PLACE1007730	14.54	9.51	13.82	11.08	11.03	10.82	7.95	12.03	8.17	*	-
PLACE1007737	26.23	17.29	21.48	12.32	15.98	9.58	13.89	18.95	16.37	*	-
PLACE1007743	15.62	16.04	17.85	8.71	10.59	7.11	6.26	11.61	4.77	**	-
PLACE1007746	42.92	48.93	37.39	28.78	37.1	35.26	18.78	10.7	15.4	**	-
PLACE1007753	14.89	18.43	15.85	13.65	30.84	24.37	8.15	10.27	7.82	**	-
PLACE1007769	10.21	11.59	12.77	11.76	15.29	7.33	6.11	7.79	6.57	**	-
PLACE1007780	5.74	2.24	8.02	2.59	3.98	3.4	2.92	4.75	4.26	*	-
PLACE1007791	7.33	7.57	11.32	6.64	9.26	5.56	5.87	21.8	5.42	*	-
PLACE1007807	10.4	6.88	6.69	5.19	10.28	4.84	5.47	16.67	5.17	*	-
PLACE1007810	21.71	5.64	13.78	5.45	11.87	20.68	5.72	5.81	5.21	*	-
PLACE1007814	23.92	18.73	27.74	15.47	17.03	13.06	8.88	14.46	7.51	*	-
PLACE1007828	9.64	11.73	14.79	8.65	13.95	8.88	3.78	6.99	9.71	*	-
PLACE1007829	21.85	26.65	22.58	10.61	17.88	12.23	16.05	20.8	14.7	*	-
PLACE1007841	12.3	14.28	18.38	8.62	14.37	8.97	13.64	10.06	9.74	*	-
PLACE1007842	7.29	10.66	10.16	4.04	8.15	4.83	12.33	8.69	6.73	*	-
PLACE1007843	8.16	7.8	8.23	4.71	8.29	4.31	4.49	6.71	2.94	*	-
PLACE1007845	14.45	11.07	14.52	8.03	11	7.42	8.69	6.18	6.07	*	-
PLACE1007846	9.62	9.87	9.68	6.82	5.35	6.88	5.9	8.42	7.28	**	-
PLACE1007848	10.98	7.82	17.21	7.21	6.58	4.31	3.05	6.37	4.05	*	-
PLACE1007852	38.45	42.72	20.94	14.02	16.83	14.13	8.88	9.78	9.37	*	-
PLACE1007856	19.55	23.44	22.4	13.38	19.15	12.08	12.88	10.77	11.27	*	-
PLACE1007849	108.08	120.7	80.75	100.84	136.45	101.33	115.22	114.23	135.63	**	-
PLACE1007871	46.64	38.13	38.79	13.91	18.93	11.35	18.19	30.79	21.75	**	-
PLACE1007877	11.11	8.45	11.03	7.02	8.84	8.93	6.39	7.32	5.13	*	-
PLACE1007878	37.18	35.47	41.1	20.44	30.52	24.13	17.29	22.96	22.69	*	-
PLACE1007881	7.08	8.99	9.43	6.21	6.72	5.42	3.38	3.24	2.37	*	-
PLACE1007885	18.54	9.97	15.34	8.52	12.53	23.99	7.37	7.86	5.04	*	-
PLACE1007897	9.52	10.47	10.76	6.15	8.22	6.75	3.33	3.79	2.67	*	-
PLACE1007908	27.14	31.91	21.03	11.57	16.17	12.14	8.02	8.92	8.32	*	-
PLACE1007922	18.27	17.29	11.92	8.56	15.72	12.01	8.76	6.61	6.69	*	-
PLACE1007946	8.9	7.88	9.69	3.7	5.72	2.73	4.49	7.73	5.93	**	-

132 / 157

2703

2704

PLACE1007950	123	99.57	80.18	78	72.07	58.93	91.95	114.45	65.58	*	-	-
PLACE1007954	8.61	7.27	7.95	5.58	7.49	5.54	3.77	4.03	2.97	**	-	-
PLACE1007955	17.84	15.71	13.68	11.51	9.3	9.77	8.99	7.04	4.12	*	-	-
PLACE1007956	11.88	9.09	7.68	6.57	3.25	6.9	4.08	4.11	3.44	*	-	-
PLACE1007958	9.03	7.99	9.04	2.85	4.9	3.85	2.85	3.75	2.07	**	-	-
PLACE1007959	15.3	15.27	14.75	7.83	11.66	10.38	7.23	6.85	6.87	*	-	-
PLACE1007960	24.54	27.08	14.83	18.49	16.9	27.44	8.5	10.51	6.83	*	-	-
PLACE1007971	14.85	11.23	10.68	4.6	5.77	4.48	5.47	4.42	6.63	**	-	-
PLACE1007990	13.72	7.91	11.82	6.44	6.58	7.05	6.02	6.88	7.23	*	-	-
PLACE1008000	8.41	4.98	5.71	4.89	6.7	6.67	3.52	3.87	3.05	*	-	-
PLACE1008002	8.38	10.22	5.77	6.22	6.73	3.91	3.28	5.34	3.1	*	-	-
PLACE1008037	7.47	6.35	4.25	2.21	4.95	4.15	2.35	1.88	2.58	*	-	-
PLACE1008044	23.79	15.95	12.5	7.05	18.88	9.73	6.75	10	8.08	*	-	-
PLACE1008045	19.99	21.06	9.24	5.88	14.67	9.04	6.52	7.55	2.12	*	-	-
PLACE1008050	15.75	26.8	10.27	8.38	25.41	7.91	16.05	14.95	13.98	*	-	-
PLACE1008092	7.87	5.27	5.41	3.04	5.53	3.46	4.86	3.3	3.17	*	-	-
PLACE1008095	18.14	13.84	10.97	4.81	15.55	7.49	3.99	8.83	6.13	*	-	-
PLACE1008105	10.75	13.42	9.63	5.45	13.95	7.7	10.03	7.98	6.35	*	-	-
PLACE1008107	6.1	7.51	5.49	4.1	6.68	4.81	2.94	3.48	3.2	**	-	-
PLACE1008111	12.94	8.4	7.21	4.85	8.37	6.82	4.58	1.47	2.41	*	-	-
PLACE1008113	49.12	32.36	38.95	30.18	28.53	18.75	11.99	9.55	15.2	**	-	-
PLACE1008122	12.52	11.61	10.25	6.69	8.27	7	4.17	3.48	4.27	**	-	-
PLACE1008129	12.19	11.22	9.29	6.85	14.98	11.14	8.9	8.48	6.51	*	-	-
PLACE1008132	21.17	14.41	14.91	9.14	12.78	11.94	8.42	8.33	8.23	*	-	-
PLACE1008137	10.44	8.69	4.43	4.58	5.28	3.39	2.11	5.99	2.53	*	-	-
PLACE1008174	12.93	8.11	5.85	5.98	7.01	8.19	6.41	5.88	5.58	*	-	-
PLACE1008177	5.92	6.81	4.83	3.39	4.65	3.28	3.39	1.53	3.81	*	-	-
PLACE1008181	12.18	13.31	12.04	4.81	8.68	5.04	4.65	2.16	4.62	**	-	-
PLACE1008195	13.39	9.3	9.62	7.44	7.78	6.35	6.1	3.44	5.91	*	-	-
PLACE1008198	12.57	11.23	9.83	6.75	6.38	4.49	7.75	4.45	5.55	**	-	-
PLACE1008201	10.88	12.64	8.33	6.13	9.68	7.85	6.3	6.05	9.18	*	-	-
PLACE1008209	10.98	9.72	8.1	4.34	4.18	5.71	10.15	7.4	9.7	**	-	-
PLACE1008226	27.87	17.88	18.73	12.29	14.5	8.3	21.53	17.31	27.74	*	-	-
PLACE1008227	19.93	14.99	13.17	9.05	11.55	7.15	13.25	11.51	13.73	*	-	-
PLACE1008231	7.21	4.7	4.12	4.69	4.49	3.28	5.49	4.08	3.39	*	-	-
PLACE1008238	12.79	18.51	15.04	5.47	8.67	11.49	8.6	7.97	4.88	*	-	-
PLACE1008244	10.53	9.75	8.28	4.31	8.05	4.21	4.37	5.27	5.5	*	-	-
PLACE1008249	15.93	17.02	8.54	6.34	10.36	6.43	17.75	4.65	8.03	*	-	-
PLACE1008266	102.89	124.96	102.56	88.83	113.4	78.29	73.83	84.82	80.68	*	-	-
PLACE1008273	14.29	12.39	14.89	6.95	10.08	11.71	9.42	8.24	5.42	*	-	-
PLACE1008275	5.26	4.99	6.79	3.64	10.8	3.9	3.08	3.54	1.54	*	-	-
PLACE1008280	8.85	8.62	8.82	4.72	8.48	4.99	3.58	6.79	2.98	*	-	-
PLACE1008282	12.89	9.41	16.7	11.11	18.66	13.55	5.47	11.9	7.88	*	-	-
PLACE1008297	12.14	13.11	10.25	7.3	9.97	7.71	3.51	6.91	7.44	*	-	-
PLACE1008323	11.47	15.79	13.39	11.12	14.28	7.92	5.84	6.87	4.98	**	-	-
PLACE1008328	4.41	7.28	5.38	6.98	9.29	6.05	3.58	4.2	4.65	*	-	-
PLACE1008315	18.59	15.1	15.12	9.88	9.1	7.98	7.41	8.94	11.54	**	-	-
PLACE1008329	9.22	7.92	9.32	5.09	12.42	9.18	10.32	5.48	5.2	*	-	-
PLACE1008330	11.04	9.84	11.21	6.66	7.9	5.74	7.19	13.58	9.01	**	-	-
PLACE1008331	9	15.63	14.68	4.1	11.97	13.67	4.88	7.64	6.55	*	-	-
PLACE1008351	12.84	22	18.28	7.81	13.2	12.68	4.26	8.16	3.26	*	-	-
PLACE1008356	13.82	13.87	16.44	11.31	18.42	14.21	4.74	8.97	9.78	*	-	-
PLACE1008359	18.61	8.87	5.75	4.5	6.41	4.58	1.38	3.42	3.83	*	-	-
PLACE1008388	15.16	18.55	16.33	11.79	13.53	6.82	4.87	5.65	5.64	**	-	-
PLACE1008389	3.81	2.08	4.52	1.83	1.71	1.5	2.12	2.77	3.88	*	-	-
PLACE1008392	11.15	8.41	8.64	4.52	7.73	5.42	7.53	11.47	5.48	*	-	-
PLACE1008394	11.83	9.88	11.91	7.27	8.44	5.81	12.55	9.98	4.42	*	-	-
PLACE1008398	23.83	18.35	28.34	11.94	10.63	17.8	18.05	34.01	14.62	*	-	-
PLACE1008401	9.05	11.72	16.92	8.35	8.24	6.49	5.59	6.98	7.72	*	-	-
PLACE1008402	8.61	13.91	16.2	7.29	15.88	9.77	4.08	3.77	5.17	*	-	-
PLACE1008405	93.35	102.54	109	58.1	65.02	51.5	64.76	83.76	83.34	**	-	-
PLACE1008408	9.15	10.93	10.45	7.65	10.58	6.32	5.13	6.5	8.18	**	-	-
PLACE1008420	4.6	3.84	4.48	2.08	3.3	3.48	3.03	3.37	1.89	*	-	-
PLACE1008424	6.77	7.89	8.59	4.05	4.03	4.27	11.48	5.37	5.37	**	-	-

133 / 157

2705

2706

PLACE1008426	9.02	4.49	7.91	3	6.94	4.35	2.2	4.69	5.02										
PLACE1008429	6.72	4.48	8.18	2.61	5.16	2.94	4.14	3.81	3.51										
PLACE1008430	14.21	10.67	13.58	6.18	13.27	13.78	7.02	13.13	6.7										
PLACE1008437	10.05	12.15	13.23	7.3	7.15	7.37	5.37	12.87	6.94	**									
PLACE1008453	9.93	7.46	12.22	2.24	6.5	6.48	4.48	4.26	3.11	*	*								
PLACE1008454	21.6	22.78	23.98	10.1	17.4	10.34	15.68	15.08	11.82	*	*								
PLACE1008455	34.62	30.32	25.55	12.02	19.33	14.29	16.61	22	23.57	*	*								
PLACE1008457	11.87	6.88	10.52	6.51	8.83	5.84	9.41	9.29	9.88	*	*								
PLACE1008465	7.78	1.83	9.85	2.4	6.54	3.49	3.88	7.05	5.33	*									
PLACE1008489	9.74	7.14	10.83	5.38	6.23	4.43	4.05	9.23	7.36	*									
PLACE1008488	7.69	9.08	16.27	2.87	27.47	11.79	3.3	10.09	3.85										
PLACE1008519	13.87	13.58	16.85	12.64	13.83	10.17	7.48	12.89	9.26										
PLACE1008524	16.01	17.28	16.34	7.24	7.92	7.96	6.91	8.21	5.65	**	**								
PLACE1008531	10.83	9.9	10.84	3.8	7.89	4.88	5.6	5.84	4.3	*	**								
PLACE1008532	31.22	27.85	29.18	13.43	17.17	10.64	20	23.65	23.08	**	**								
PLACE1008533	16.83	14.02	15.82	5.16	9.58	6.45	8.04	10.1	5.87	**	**								
PLACE1008542	11.05	6.89	12.38	7.82	8.01	5.6	3.52	4.43	4.89	*									
PLACE1008549	9.08	8.32	6.58	2.78	4.31	2.64	3.18	4.98	1.09	**	*								
PLACE1008560	9.22	6.82	8.11	4.41	4.7	3.88	2.93	3	1.88	**	**								
PLACE1008587	12.48	12.8	11.19	3	4.87	3.75	6.88	4.77	3.6	**	**								
PLACE1008588	11.23	14.36	12.88	7.02	10.87	9.2	4.7	4.88	4.71	*	**								
PLACE1008589	26.78	24.83	18.12	14.18	21.2	18.33	7.27	6.81	8.32	**									
PLACE1008594	12.08	11.9	8.94	4.22	7.88	8.89	3.89	8.17	12.55										
PLACE1008595	31.41	26.84	21.37	12.16	18.6	14.29	13.28	12.03	8.85	*	**								
PLACE1008603	68.44	68.88	85.85	77.22	187.17	81.02	46.89	50.43	38.78	*									
PLACE1008621	8.51	5.24	9.31	3.02	5.85	4.1	5.45	4.49	4.31										
PLACE1008625	5.93	4.92	6	3.64	3.65	3.18	1.59	3.39	1.87	**	**								
PLACE1008628	7.17	7.14	8.41	4.47	6	4.22	1.34	2.51	2.04	*	**								
PLACE1008627	18.53	17.04	15.47	5.85	18.41	6.49	6.26	8.1	3.78	**	**								
PLACE1008629	17.85	11.92	13.81	7.66	28.73	20.07	19.18	21.96	10.11										
PLACE1008630	10.34	9.61	9.25	3.39	5.31	3.93	4.34	4.78	3.98	**	**								
PLACE1008643	13.74	8.08	11.06	4.37	9.48	6.43	8.52	10.48	6.53										
PLACE1008650	8.08	8.41	8.78	5.58	6.65	5.55	2.49	3.07	2.46	**									
PLACE1008657	20.07	15.69	13.65	10.15	11.39	11.63	13.41	12.65	11.83	*									
PLACE1008664	16.27	13.14	9.24	7.33	8.26	11.12	3.83	4.48	3.87	*									
PLACE1008697	19.24	12.55	11.48	5.9	7.48	8.22	10.23	10.78	7.26	*									
PLACE1008696	15.27	18.92	13.52	8.86	14.35	6	9.09	6.63	6.3	**									
PLACE1008713	12.07	13.35	10.87	4.79	8.42	4.38	3.79	3.86	5.13	*	**								
PLACE1008716	9.95	8.16	6.63	2.27	4.4	4.51	4.19	6.11	4.54	*	*								
PLACE1008722	23.08	17.9	16.82	6.63	6.45	7.4	2.92	4.69	5.05	**	**								
PLACE1008738	22.32	18.29	21.17	13.47	18.87	5.51	15.83	18.57	23.28										
PLACE1008742	84.96	67.74	76.79	63.3	63.65	55.26	60.09	57.16	55.95	*	*								
PLACE1008744	16.51	21.18	12.37	7.32	26.18	7.33	4.32	8.49	4.3	*									
PLACE1008748	11.7	8.68	7.89	3.6	3.19	1.95	5.93	2.92	4.48	**	*								
PLACE1008757	11.88	10.89	13.3	7.9	13.9	4.82	5.45	7.52	5.79	**									
PLACE1008765	12.68	9.83	17.07	7.19	5.82	5.22	3.78	3.59	4.47	*	*								
PLACE1008785	10.18	7.84	9.9	4.81	6.58	4.55	5.8	6.94	7.6	*									
PLACE1008790	13.83	10.34	7.69	5.81	9.25	7.49	3.5	9.44	5.2										
PLACE1008798	28.65	17.87	28.15	8.38	18.4	13.81	20.64	7.92	18.2										
PLACE1008807	7.38	8.08	6.82	3.5	7.58	4.57	3.41	3.28	4.62	**									
PLACE1008808	14.69	14.79	11.52	7.99	7.57	5.08	6.28	4.51	3.87	**	**								
PLACE1008813	11.75	10.29	9.89	7.89	7.9	5.45	5.87	3.03	5.81	*	**								
PLACE1008838	10.67	11.89	10.55	8.13	8.4	5.79	6.93	5.51	7.49	**	**								
PLACE1008851	17.38	11.13	11	56.43	9.1	8.6	16.81	8.29	14.77										
PLACE1008854	6.2	6.11	5.28	5.3	4.42	2.5	6.46	5.33	5.1										
PLACE1008884	19.27	13.51	11	10.42	10.96	6.48	14.43	10.21	14.23										
PLACE1008887	23.08	16	11.22	6.2	8.84	4.88	5.54	7.61	6.44	*									
PLACE1008876	176.15	214.21	204.78	126.27	307.85	120.08	159.59	183.82	275.73										
PLACE1008887	18.24	8.9	12.49	4.83	6.78	7.78	5.17	4.7	6.53	*	*								
PLACE1008902	13.44	7.31	7.6	4.55	6.21	3.19	5.49	3.64	7.36										
PLACE1008911	28.57	23.29	15.12	14.15	10.78	7.02	9.39	4.72	9.35	*	*								
PLACE1008917	7.8	5.68	7.21	2.11	3.42	3.23	3.5	4.03	3.28	**	**								
PLACE1008920	3.68	3.98	2.25	1.85	2.38	2.78	2.45	2.52	1.93										
PLACE1008925	6.03	2.49	3.19	2.51	4.89	2.3	3.22	3.16	2.78										

134 / 157

2707

2708

PLACE1008930	19.58	15.04	12.13	4.18	13.75	5.17	6.22	5.55	4.1	*	-
PLACE1008934	18.24	5.52	7.38	4.49	6.52	4.37	6.49	7.17	5.19	*	-
PLACE1008941	7.73	5.12	5.84	2.98	5.21	1.59	3.04	3.68	3.45	**	-
PLACE1008947	14.65	13.48	10.43	5.14	6.75	4.89	11.51	10.32	9.84	**	-
PLACE1008954	13.25	12.98	9.34	5.45	4.62	4.47	6.01	1.98	3.16	**	-
PLACE1008955	7.07	4.44	3.91	3.08	5.06	1.59	3.53	4.2	7.08	*	-
PLACE1008994	3.13	3.32	2.48	1.43	1.88	1.44	1.4	0.58	1.95	**	-
PLACE1009020	10.8	10.64	11.79	9.97	10.93	10.07	6.16	7.97	6.48	**	-
PLACE1009027	3.76	1.41	3.27	1.39	2.64	1.03	1.54	3.8	0.71	*	-
PLACE1009039	4.77	3.08	3.94	1.85	3.87	1.57	1.09	2.8	1.86	*	-
PLACE1009045	8.62	6.04	9.52	3.29	4.82	2.76	2.8	7.24	2.61	*	-
PLACE1009048	2.08	2.51	4.83	1.04	3.38	1.38	0.62	1.42	2.18	*	-
PLACE1009050	3.79	3.8	4.82	1.78	8.12	1.71	2.85	3.16	1.84	*	-
PLACE1009080	19.41	22.53	18.93	15.31	14.46	12.87	9.32	9.83	14.15	*	-
PLACE1009087	9.68	7.46	10.13	4.56	10.58	6.42	9.08	6.46	12.36	*	-
PLACE1009071	20.58	20.38	20.62	12.02	12.73	10.86	11.69	13.76	13.78	**	-
PLACE1009090	9.2	7.98	10.33	4.26	8.7	11	5.63	8.05	9.63	*	-
PLACE1009091	83.22	75.58	81.63	64.89	72.29	69.89	50.11	57.74	52.35	*	-
PLACE1009094	10.97	9.99	9.32	7.26	14.47	8.52	5.42	12.88	9.03	*	-
PLACE1009098	8.3	10.72	9.62	5.38	7.81	7.04	3.55	4.21	4.97	*	-
PLACE1009110	7.1	4.33	5.05	4.77	6.82	3.94	3.19	5.04	4.48	*	-
PLACE1009111	4.96	7.16	3.67	6.5	7.91	3.83	4.35	3.44	4.4	*	-
PLACE1009112	16.47	13.6	10.85	8.18	11.25	17.03	7.41	8	12.78	*	-
PLACE1009130	13.28	8.46	18.05	9.81	8.47	7.77	7.71	8.47	5.51	*	-
PLACE1009150	9.24	9.08	10.58	6.69	8.31	6.53	5.87	7.78	4.65	*	-
PLACE1009155	16.03	18.94	19.94	9.19	13.17	10.27	14.29	28.01	20.9	*	-
PLACE1009158	11.31	7.26	10.08	8.16	8.65	7.83	4.47	5.84	5.53	*	-
PLACE1009165	8.72	5.63	8.02	5.97	6.75	3.66	4.53	5.03	5.29	*	-
PLACE1009172	12.12	10.58	13.44	7.25	11.43	5.18	4.32	7.13	6.22	**	-
PLACE1009174	15.11	18.12	12.65	18.74	28.86	11.28	7.87	11.41	11.21	*	-
PLACE1009183	26.21	21.05	18.45	14.71	13.12	9.63	13.01	10.13	17.14	*	-
PLACE1009185	18.33	12.69	14.24	9.34	9.68	8.31	3.27	14.09	4.74	**	-
PLACE1009188	10.58	9.79	11.18	7.65	14.08	8.19	4	6.56	7.14	*	-
PLACE1009189	13.22	10.75	11.8	7.38	9.71	7.63	10.55	12.67	8.91	*	-
PLACE1009200	19.33	23.87	22.85	10.45	11.73	10.18	10.96	16.99	17.03	**	-
PLACE1009217	14.13	24.09	18.64	11.71	19.57	10.58	3.82	4.84	6.12	*	-
PLACE1009230	27.31	26.63	32.64	10.81	26.12	18.34	16.83	20.7	24.31	*	-
PLACE1009236	9.03	10.62	12.67	7.18	9.87	8.71	6.77	8.24	7.63	*	-
PLACE1009246	21.48	34.44	20.89	9.28	28.22	18.2	9.84	35.09	13.46	*	-
PLACE1009255	18.88	26.11	26.51	7.08	18.04	8.4	9.34	41.89	9.73	*	-
PLACE1009279	13.38	8.53	11.2	5.1	7.71	8.09	11.6	8.18	8.91	*	-
PLACE1009285	49.41	26.44	34.52	26.24	34.87	27.56	30.84	25.04	29.21	*	-
PLACE1009308	8.89	11.34	12.22	3.57	6.53	4.13	6.2	4.88	5.8	**	-
PLACE1009319	11.98	14.82	14.36	7.73	14.05	5.85	5.29	8.69	4.23	*	-
PLACE1009328	12.86	13.58	16.59	6.43	18.45	5.88	7.1	8.23	11.41	*	-
PLACE1009335	8.34	6.5	7.11	3.99	9.88	4.55	4.35	7.62	2.32	*	-
PLACE1009338	14.56	12.99	17.79	6.9	13.74	10.82	6.69	5.83	8.27	**	-
PLACE1009344	6.92	5.13	5.88	3.02	11.98	4.73	3.18	3.88	2.68	*	-
PLACE1009355	23.17	22.99	28.45	20.27	18.4	17.42	10.13	10.79	9.83	*	-
PLACE1009368	8.63	5.08	8.32	3.35	3.68	2.95	3.35	6.92	2.66	*	-
PLACE1009375	8.35	5.94	11.71	3.65	4.61	3.52	3.31	4.45	3.97	*	-
PLACE1009388	18.28	22.47	42.89	8.1	17.38	17.02	11.33	10.89	17.6	*	-
PLACE1009398	16.02	15.66	15.45	5.65	12.56	7.13	6.61	5.4	4.18	*	-
PLACE1009404	17.98	10.01	28.67	4.73	28.94	16.4	9.28	9.88	5.15	*	-
PLACE1009410	13.78	15.39	9.6	5.62	5.24	5.56	5.28	4.84	5.63	*	-
PLACE1009417	18.73	9.41	10.65	12.79	9.15	6.2	6.71	6.82	7.05	*	-
PLACE1009424	9.71	6.5	9.08	4.75	4.94	4.28	3.44	4.18	4.16	*	-
PLACE1009434	21.73	12.22	10.01	6.12	7.87	5.95	3.69	5.83	2.87	*	-
PLACE1009443	7.33	14.79	8.94	2.98	3.24	2.54	4.85	2.08	4.82	*	-
PLACE1009444	13.13	14.77	17.33	7.6	10.32	8.01	6.52	5.24	6.76	*	-
PLACE1009459	8.33	7.91	9.2	3.33	5.72	4.79	3.49	3.56	3.33	**	-
PLACE1009460	9.94	14.6	11.32	7.7	8.3	7.62	5.02	3.81	2.95	*	-
PLACE1009468	12.43	12.85	9.11	5.4	5.98	4.76	4.66	5.65	8.17	**	-
PLACE1009476	7.02	4.9	5.38	3	3.07	3.32	3.74	3.78	3.51	*	-

135 / 157

2709

2710

PLACE1009477	13.59	11.11	12.8	10.3	8.83	6.89	8.76	11.16	10.89				
PLACE1009493	5.63	4.68	4.85	3.04	3.33	2.25	3.05	2.29	1.49	**	**	-	-
PLACE1009502	11.43	9.62	8.39	5.92	5.81	5.8	4.63	7.61	3.19	*	*	-	-
PLACE1009524	7.58	7.28	7.63	1.87	3.68	2.97	2.68	3.37	2.45	**	**	-	-
PLACE1009527	18.34	12.36	8.71	6.88	11.49	9.95	6.88	5.14	3.52	*	*	-	-
PLACE1009531	26.38	45.76	31.03	26.02	25.83	22.82	19.07	17.64	15.03				
PLACE1009535	12.3	11.08	8.3	3.83	8.2	5.84	3.35	3.71	4.06	*	**	-	-
PLACE1009538	8.63	10.23	7.02	4.34	7.58	5.81	5.39	6.61	6.43				
PLACE1009540	37.43	26.63	27.99	14.5	25.74	28.63	21	11.9	8.28	*	*	-	-
PLACE1009542	12	5.45	5.7	3.62	4.51	6	3.18	3.89	2.87				
PLACE1009546	12.67	10.42	8.28	3.72	6.38	5.88	4.25	10.8	3.68	*	**	-	-
PLACE1009556	7.91	8.37	8.72	5.1	4.62	3.16	3.16	2.46	2.96	*	**	-	-
PLACE1009569	11.99	15.44	10.67	4.78	13.77	8.59	5.94	5.33	6.24	**	**	-	-
PLACE1009571	7.82	9.37	5.59	3.8	7.13	3.69	5.84	2.83	4.48				
PLACE1009573	22.09	15.06	12.31	9.78	17.27	9.7	8.26	8.91	7.87	*	*	-	-
PLACE1009576	12.53	9.49	8.65	4.21	8.31	4.13	5.49	5.78	4.15	*	*	-	-
PLACE1009580	9.88	9.33	7.56	5.13	7.69	4.5	8.75	3.58	4.85				
PLACE1009581	12.95	12.1	8.15	7.19	5.84	4.99	3.45	2.48	3.63	*	**	-	-
PLACE1009587	13.3	7.18	8.28	7.05	8.26	5.41	3.85	4.9	4.28				
PLACE1009593	18.08	14.97	12.54	14.73	10.78	9.76	5.8	5.68	6.75	**	**	-	-
PLACE1009595	24.28	22.3	15.19	12.22	14.78	12.77	19.19	14	13.61				
PLACE1009596	8.31	5.39	3.71	3.49	5.52	2.24	4.95	2.48	3.3	*	*	-	-
PLACE1009600	19.52	17.07	12.01	6.59	8.79	3.99	8.49	7.24	8.2	*	*	-	-
PLACE1009604	19.98	10.38	8.88	4.18	6.3	6.68	3.83	5.54	4.55				
PLACE1009607	17.2	18	14.19	8.25	8.63	7.56	19.64	17.82	14.33	**	**	-	-
PLACE1009613	8.31	8.44	8.08	5.1	6.18	4.22	4.39	2.55	3.96	*	*	-	-
PLACE1009621	18.02	17.68	15.95	7.12	14.78	8.85	8.46	6.45	12.83	*	*	-	-
PLACE1009622	18.83	8.51	8.44	5	5.84	4.48	4.98	3.18	4.58				
PLACE1009624	28.04	18.02	15.23	8.31	9.71	5.05	7.94	5.67	7.85	*	**	-	-
PLACE1009637	8.56	7.93	7.76	3.81	7.55	4.55	3.93	3.27	4.29	**	**	-	-
PLACE1009639	14.9	6.92	2.67	7.62	5.31	8.34	6.06	5.89	7.84				
PLACE1009654	23.03	16.88	13.87	7.6	11.88	8.68	7.4	8.09	10.22	*	*	-	-
PLACE1009658	17.62	16.71	15.38	8.59	15.54	12.83	6.78	8.01	7.95	**	**	-	-
PLACE1009665	13.55	12.11	8.08	5.71	10.68	10.87	8.53	10.59	6.01				
PLACE1009669	10.91	11.58	11.44	5.71	8.38	5.23	8.32	8.02	8.86	**	**	-	-
PLACE1009670	10	4.03	5.22	3.83	4.76	8.45	5.57	4.18	5.29				
PLACE1009708	13.41	10.89	7.63	11.97	12.83	3.57	6.84	3.75	9.58				
PLACE1009721	16.78	10.82	9.31	5.16	8.52	4.42	9.11	3.24	6.91	*	*	-	-
PLACE1009731	5.6	6.57	11.05	5.5	8.58	5.78	5.45	4.8	9.13				
PLACE1009735	9.43	10.36	12.52	8.67	8.77	6.91	8.15	10.58	12.7				
PLACE1009737	8.38	8.02	10.88	5.74	17.02	9.89	4.88	11.47	4.68				
PLACE1009741	8.67	7.59	11.34	6.59	7.8	4.16	2.83	4.89	4.5	*	*	-	-
PLACE1009752	9.51	12.78	18.39	7.48	12.57	11.15	4.36	9.17	8.85				
PLACE1009763	16.81	19.39	15.73	12.5	15.24	13.69	4.1	8.02	6.41	**	**	-	-
PLACE1009766	7.54	8.76	7.16	6.98	11.81	6.18	9.17	7.21	6.61				
PLACE1009772	12.62	18.28	13.48	18.58	26.84	15.97	7.94	11.57	13.61				
PLACE1009782	7.66	6.95	7.89	4.24	8.2	9.33	6.26	4.41	4.35	*	*	-	-
PLACE1009794	8.71	9.98	15.31	6.87	7.94	5.84	5.81	8.88	9.35				
PLACE1009798	15.7	11.58	19.23	8.55	9.28	7.81	5.78	14.83	6.88	*	*	-	-
PLACE1009845	5.68	8.07	10.29	5.79	8.81	8.4	3.27	3.21	4.35	*	*	-	-
PLACE1009849	4.28	8.11	7.54	5.28	8.21	5.62	3.81	4.56	5.57				
PLACE1009857	5.97	7.82	9.09	4.47	4.42	3.68	4.36	2.37	3.88	*	*	-	-
PLACE1009861	21.52	19.65	20.31	10.08	12.87	8.67	13.97	16.84	18.62	**	**	-	-
PLACE1009872	225.97	496.72	352.07	405.22	548.08	335.22	583.26	819.73	801.38	*	*	-	-
PLACE1009877	93.23	95.23	121.84	37.51	28.89	31.48	33.83	34.28	50.76	**	**	-	-
PLACE1009879	8.31	8.4	8.23	4.6	3.04	4.45	5.4	6.04	6.14	**	**	-	-
PLACE1009885	6.21	5.71	8.04	3.1	4.43	3.64	3.76	3.74	3.91	**	**	-	-
PLACE1009888	16.61	15.12	18.88	7.12	11.3	10.4	7.67	15.82	6.68	*	*	-	-
PLACE1009908	10.69	13.22	16.94	7.9	15.1	13.2	5.89	21.59	11.16				
PLACE1009919	18.85	20.8	15.18	25.14	44.25	16.53	8.38	10.08	17.42				
PLACE1009921	12.5	11.43	13.89	6.9	11.45	10.84	6.74	7.53	8.61	**	**	-	-
PLACE1009923	37.38	53.14	35.03	28.71	68.72	45.4	23.4	36.13	28.69				
PLACE1009924	28.03	18.87	21.48	6.84	14.99	12.69	28.73	40.32	10.81				
PLACE1009925	18.23	25.64	14.3	6.85	14.35	14.82	6.32	35.83	7.94				

136 / 157

2711

2712

PLACE1009931	29.02	24.07	26.64	12.24	15.6	14.51	8.57	9.79	10.88	**	**	-	-
PLACE1009935	8.88	7.44	9.7	2.78	11.02	5.42	14.95	3.43	6.35	*	*	-	-
PLACE1009947	8.29	6.77	9.83	4.26	4.43	3.14	2.04	2.93	4.01	*	*	-	-
PLACE1009981	7.86	10.21	16.93	5.24	6.64	4.49	4.58	6.45	5	*	*	-	-
PLACE1009971	7.79	7.18	10.28	3.38	7.18	3.34	3.67	4.69	3.08	*	*	-	-
PLACE1009982	20.35	18.31	15.25	6.6	11.54	11.1	7.87	5.59	13.47	*	*	-	-
PLACE1009992	12.32	8.01	11.62	6.39	8	6.72	5.24	5.76	6.38	*	*	-	-
PLACE1009995	21.27	22.11	27.33	13.99	19.4	11.81	8.12	18.7	12.66	*	*	-	-
PLACE1009997	23.76	8.03	15.68	6.79	12.9	5.63	10.63	9.44	8.19	*	*	-	-
PLACE1010002	14.66	6.35	12.19	4.68	8.08	5.35	5.32	14.28	7.75	*	*	-	-
PLACE1010011	15.51	14.11	17.68	6.19	7.34	6.29	9.9	6.46	10.43	**	*	-	-
PLACE1010012	6.16	5.13	12.68	2.37	2.99	1.58	3.53	4.24	2.88	*	*	-	-
PLACE1010021	15.95	20.29	19.12	7.11	14.51	9.31	5.22	8.04	6.15	*	**	-	-
PLACE1010023	27.49	31.11	21.32	17.41	29.7	20.44	16.49	7.28	6.33	*	*	-	-
PLACE1010031	20.87	15.68	32.54	10.16	16.34	7.15	9.85	6.36	10.28	*	*	-	-
PLACE1010039	20.5	8.82	10.22	3.99	6.51	3.25	2.91	6.23	3.6	*	*	-	-
PLACE1010045	11.17	9.8	13.16	5.75	8.43	6.1	5.11	5.8	6.51	*	**	-	-
PLACE1010053	7.04	5.22	9.04	6.47	4.19	2.39	2.32	2.12	1.18	*	*	-	-
PLACE1010060	18.9	18.89	15.21	7.4	6.50	8.37	12.03	13.21	10.77	**	*	-	-
PLACE1010069	10.06	8.9	13.66	5.33	6.93	4.79	2.88	4.42	3.79	*	*	-	-
PLACE1010070	8.15	7.09	9.72	2.34	9.93	4.09	3.65	3.87	2.87	**	*	-	-
PLACE1010074	63.58	56.74	39.11	24.21	42.72	26.72	25.32	15.29	17.22	*	*	-	-
PLACE1010076	60.58	44.35	88.45	21.22	23.2	19.4	13.41	12.75	17.59	**	**	-	-
PLACE1010078	13.76	11.23	14.35	6.69	10.2	6.63	4.38	4.87	4.45	*	**	-	-
PLACE1010081	27.34	19.16	21.67	12.55	15.05	11.39	11.73	7.47	11.77	*	*	-	-
PLACE1010083	7.1	7.48	5.64	1.97	2.84	2.62	1.69	2.61	1.96	**	**	-	-
PLACE1010089	11.95	9.85	9.57	5.6	5.81	4.38	4.84	2.48	4.43	**	**	-	-
PLACE1010095	14.85	29.76	16.35	5.81	14.35	9.63	7.28	8.67	5.49	*	*	-	-
PLACE1010102	22.29	34.08	23.15	9.26	15.03	13.51	7.63	8.06	7.81	*	**	-	-
PLACE1010105	18.16	20.35	15.89	6.52	12.73	7.6	4.02	4.82	4.1	*	**	-	-
PLACE1010108	19.44	22.57	15.73	8.29	8.94	12.84	10.96	10.43	10.44	*	*	-	-
PLACE1010130	6.82	6.56	4.7	3.25	3.39	2.18	3.26	3.47	3.87	*	*	-	-
PLACE1010132	20.18	25.39	20.58	20.18	19.1	12.29	12.71	11.47	12.5	**	**	-	-
PLACE1010134	10.35	11.42	9.66	6.05	6.05	5.55	6.72	4.12	2.88	**	**	-	-
PLACE1010139	74.68	85.11	79.85	37.26	51.16	43.94	15.88	26.17	18.5	**	**	-	-
PLACE1010140	10.48	7.43	9.71	13.14	7.54	4.66	3.27	4.13	7.63	*	*	-	-
PLACE1010155	9.79	7.58	7.83	3.3	5.89	2.96	2.11	2.28	2.53	*	**	-	-
PLACE1010156	32.69	26.06	24.61	14.7	18.09	11.09	7.29	10.58	14.48	*	**	-	-
PLACE1010181	8.48	13.16	10.69	5.15	8.21	6.03	4.47	3.31	5.48	*	*	-	-
PLACE1010181	8.66	8.08	8.51	2.62	6.54	5.02	3.74	4.12	2.93	**	*	-	-
PLACE1010194	8.57	7.46	8.67	3.93	4.98	3.84	2.85	2.08	2.37	**	**	-	-
PLACE1010202	8.39	6.76	12.45	8.31	10.9	5.79	3.58	3.42	2.04	*	*	-	-
PLACE1010231	12.97	10.31	14.49	7.98	15.61	7.14	8.93	9.42	8.37	*	*	-	-
PLACE1010235	12.62	15	11.24	4.59	11.54	4.53	6.11	5.14	4.78	**	*	-	-
PLACE1010237	5.04	3.77	4.4	2.18	2.37	2.25	2.72	0.64	1.32	**	*	-	-
PLACE1010251	9.48	4.47	6.29	5.09	4.79	10.14	3.78	5.92	3.68	*	*	-	-
PLACE1010261	8.26	6.41	4.7	3.42	2.88	2.61	4.52	5.73	4.64	*	*	-	-
PLACE1010270	7.53	8.07	8.36	3.29	6.44	3.19	3.85	4.95	3.8	**	*	-	-
PLACE1010273	8.97	10.55	5.7	2.72	5.68	3.04	2.78	2.04	3.83	*	*	-	-
PLACE1010274	20.97	18.72	29.58	10.68	16.29	14.18	6.73	5.73	6.16	**	*	-	-
PLACE1010277	15.29	14.99	14.3	8.87	13.93	8.53	11.24	4.25	5.68	*	*	-	-
PLACE1010283	16.94	20.61	17.07	9.41	11.83	8.24	17.5	8.1	13.23	**	*	-	-
PLACE1010297	38.08	33.94	34.95	17.91	31.89	16.63	31.6	20.1	25.32	*	*	-	-
PLACE1010900	21.55	15.58	11.87	9.15	8.78	7.88	3.82	5.36	5.08	*	*	-	-
PLACE1010310	323.02	293.14	231.29	170.38	221.98	136.69	214.58	235.35	207.22	*	*	-	-
PLACE1010321	10.7	10.58	7.81	4.88	7.93	6.91	5.6	6.6	6.56	*	*	-	-
PLACE1010324	6.25	5.69	6.54	2.88	4.51	2.8	4.43	3.34	2.72	*	**	-	-
PLACE1010329	14.25	11.37	11.93	5.93	10.64	4.89	9.64	9.84	9.32	*	*	-	-
PLACE1010330	12.28	14.21	10.49	11.84	10.18	7.63	12	7.94	17.88	*	*	-	-
PLACE1010335	27.7	52.68	38.23	18.39	9.7	13.5	16.61	15.28	28.61	*	*	-	-
PLACE1010341	6.44	4.91	4.72	2	3.7	1.15	4.14	2.93	4.21	*	*	-	-
PLACE1010342	2.35	2.93	2.85	2.18	1.61	1.84	1.14	1.19	0.83	*	**	-	-
PLACE1010346	10.42	6.88	7.23	4.08	5.74	2.77	5.84	5.92	5.88	*	*	-	-
PLACE1010362	13.25	8.57	9.24	6.56	7.08	3.39	11.5	13.76	14.05	*	*	-	-

137 / 157

2713

2714

PLACE1010384	11.29	7.43	7.22	2.26	3.51	2.16	4.95	3.55	4.16	*	*	-	-
PLACE1010388	10.78	8.61	7.03	3.90	9.56	5.19	7.58	7.15	5.73	*	*	-	-
PLACE1010379	30.2	28.61	24.17	14.82	22.58	10.40	24.63	20.09	20.87	*	*	-	-
PLACE1010383	18.56	18.15	18.04	10	15.54	3.52	27.13	12.41	12.03	*	*	-	-
PLACE1010385	2.48	1.73	1.18	0.83	2.33	0.48	2.65	1.19	2.08	*	*	-	-
PLACE1010389	8.48	7.51	8.42	3.76	4.47	5.47	3.96	4.51	2.91	**	**	-	-
PLACE1010401	3.24	0.96	4.6	2.29	2.49	1.63	2.81	2.31	1.6	*	*	-	-
PLACE1010410	22.91	15.39	21.28	8.31	20.95	12.79	12.79	10.9	11.3	*	*	-	-
PLACE1010418	18.78	19.01	17.17	5.75	13.45	6.67	11.25	18.37	11.11	*	*	-	-
PLACE1010425	10.58	10.75	18.36	4.2	4.88	3.19	8.87	11.17	8.75	*	*	-	-
PLACE1010443	12.48	14.66	18.77	5.6	14.99	6.09	8.79	8.23	11.76	*	*	-	-
PLACE1010445	36.17	45.7	37.43	17.66	20.28	18.29	39.23	32.45	41.58	**	**	-	-
PLACE1010481	50.4	48.95	65.37	22.92	28.42	21.44	27.71	30.53	27.4	**	**	-	-
PLACE1010482	127.4	82.31	84.53	104.64	120.51	109.85	62.40	50.87	49.87	*	*	-	-
PLACE1010491	71.28	72.3	73.41	43.51	47.95	35.09	68.63	56.51	74.84	**	**	-	-
PLACE1010492	20.32	20.74	19.72	20.62	27.04	23.27	8.07	12.55	10.15	**	**	-	-
PLACE1010509	11.2	13.07	17.63	8.18	12.58	8.24	5.79	8.34	7.88	*	*	-	-
PLACE1010518	9.76	12.31	18.67	8.17	8.57	7.5	8.04	6.33	6.73	*	*	-	-
PLACE1010522	8.32	9.35	9.63	6.71	11.42	6.63	6.37	6.88	7.65	*	*	-	-
PLACE1010528	13.09	22.15	14.64	11.31	17.06	8.68	7.61	9.45	9.29	*	*	-	-
PLACE1010547	26.79	34.33	38.34	12.39	16.81	11.19	21.21	24.3	23.05	**	**	-	-
PLACE1010560	10.15	9.34	9.68	6.09	4.54	5.89	6.02	6.35	6.13	**	**	-	-
PLACE1010582	4.79	4.38	12.01	8.8	8.65	6.38	8.41	7.48	4.4	*	*	-	-
PLACE1010575	74.54	67.98	59.08	46.33	48.92	48.24	56.27	64.49	60.31	*	*	-	-
PLACE1010580	38.79	45.7	48.75	22.24	30	17.17	31.95	38.27	36.2	*	*	-	-
PLACE1010598	28.35	28.84	32.71	12.61	15.12	14.18	19.83	24.34	23.16	**	**	-	-
PLACE1010626	31.78	30.37	40.62	15.48	19.15	14.23	22.77	25.83	28.81	**	**	-	-
PLACE1010618	16.39	26.11	18.04	12.88	17.78	7.33	3.49	3.67	4.82	**	**	-	-
PLACE1010622	9.08	4.75	12.08	2.52	4.26	2.63	11.64	11.12	6.28	*	*	-	-
PLACE1010624	292.79	228.16	204.46	212.8	278.86	210.51	168.11	171.99	155.01	*	*	-	-
PLACE1010628	48.68	44.83	61.05	27.25	26.82	37.41	15.79	18.04	21	*	*	-	-
PLACE1010629	19.03	17.43	20.71	12.12	18.38	15.36	14.93	10.66	11.67	*	*	-	-
PLACE1010630	15.62	16.61	28.46	8.82	19.7	10.88	11.96	14.1	11.42	*	*	-	-
PLACE1010631	22.99	34.46	25.64	20.07	21.82	17.69	19.23	10.89	18.98	*	*	-	-
PLACE1010651	8	10.03	11.34	6.4	9.54	5.75	4.35	5.43	5.06	**	**	-	-
PLACE1010661	13.76	16.34	32.78	7.67	13.64	27.31	8.28	10.55	7.26	*	*	-	-
PLACE1010682	28.68	22.75	30.53	8.67	12.27	9.85	25.45	28.04	24.56	**	**	-	-
PLACE1010686	37.33	37.7	40.85	22.82	25.66	20.72	38.78	44.21	37.48	**	**	-	-
PLACE1010702	12.3	6.26	13.51	6.5	7.73	5.88	8.16	17.08	6.48	*	*	-	-
PLACE1010709	70.65	75.49	65.95	55.08	47.8	67.04	60.07	52.69	55.39	**	**	-	-
PLACE1010713	80.41	68.48	77.71	35.46	48.82	48.26	28.63	32.14	23.64	**	**	-	-
PLACE1010714	14.85	12.78	15.24	4.19	7.98	4.28	3.33	8.97	5.69	**	*	-	-
PLACE1010716	10.07	15.73	17.8	4.88	22.04	9.1	5.99	13.73	3.7	*	*	-	-
PLACE1010717	18.27	25.02	15.64	13.63	18.64	10.49	10.76	15.18	3.55	*	*	-	-
PLACE1010720	27.49	32.65	34.2	14.65	18.34	8.25	20.45	22.85	22.74	**	*	-	-
PLACE1010739	32.05	33.82	24.71	15.33	21.42	16.16	17.5	28.68	18.95	*	*	-	-
PLACE1010743	44.76	28.81	34.77	15.52	24.11	19.41	14.02	9.67	5.8	*	**	-	-
PLACE1010752	26.13	19.46	22.02	11.29	9.49	11.37	16.27	24.35	13.97	**	**	-	-
PLACE1010761	16.13	12.05	32.55	16.98	9.45	8.78	11.14	13.99	12.89	*	*	-	-
PLACE1010771	28.91	75.96	37.99	25.83	90.51	39.2	31.02	18.86	25.47	**	**	-	-
PLACE1010784	98.45	116.91	93.11	48.56	63.14	52.26	30.32	31.37	33.38	**	**	-	-
PLACE1010786	18.19	15.53	16.87	8.23	15.52	9.85	7	6.49	9.87	**	**	-	-
PLACE1010789	48.25	32.28	40.6	14.8	19.17	10.09	44.84	47.22	35.88	**	**	-	-
PLACE1010800	13.97	9.78	14.67	8.91	5.6	5.82	4.05	5.43	4.76	*	**	-	-
PLACE1010802	13.35	16.74	27.18	16.21	23.88	4.91	12.32	5.78	37.65	*	*	-	-
PLACE1010811	26.8	21.58	31.28	13.09	18.01	12.69	11.33	11.86	10.25	*	**	-	-
PLACE1010813	18.21	6.96	9.67	4.84	4.38	4.87	3.92	2.83	2.64	*	**	-	-
PLACE1010827	10.79	10.81	12.05	5.43	7.48	5.67	6.85	5.02	3.78	**	**	-	-
PLACE1010833	9.18	9.88	8.82	3.14	6.94	3.42	1.97	2.07	1.97	*	**	-	-
PLACE1010839	24.14	30.08	20.18	11.7	20	12.82	18.21	11.13	15.42	*	*	-	-
PLACE1010856	16.47	10.08	17.88	6.16	8.58	4.26	11.23	8.32	12.26	*	*	-	-
PLACE1010857	32.67	31.48	18.71	21.99	18.34	15.41	16.23	15.82	15.61	*	**	-	-
PLACE1010870	5.83	4.92	4.28	3.87	3.55	2.92	2.91	2.04	1.94	*	**	-	-
PLACE1010877	26.07	20.6	20.89	14.13	8.42	9.71	13.42	12.71	7.25	**	*	-	-

138 / 157

[0962]

[表491]

2715

2716

PLACE1010882	17.08	13.64	8.42	6.17	5.89	5.82	7.13	5.01	5.37	*	-
PLACE1010891	27.43	20.32	21.15	12.48	14.41	7.48	15.28	15.07	24.71	*	-
PLACE1010896	13.62	18.23	16.16	6.92	8.03	6.78	3.64	4.82	4.08	**	**
PLACE1010900	88.8	75.85	67.62	55.88	59.03	45.47	67.12	53.92	39.14	*	-
PLACE1010916	104.52	71.53	76.5	35.13	35.18	33.1	29.84	23.73	27.17	**	**
PLACE1010917	14.21	14.67	11.27	7.82	5.59	7.41	7.89	7.08	8.78	**	**
PLACE1010924	5.86	2.7	4.5	2.89	4.47	3.08	2.47	3.1	1.31	*	-
PLACE1010925	4.94	4.55	4.99	1.88	2.53	1.67	3.09	1.69	1.78	**	**
PLACE1010926	13.94	11.18	11.63	5.87	7.64	6.22	5.6	4.23	4.29	**	**
PLACE1010942	207.77	196.21	175.48	147.15	130.24	102.36	197.42	120.16	123.73	*	-
PLACE1010943	31.5	28.2	22.88	12.49	22.18	9	15.12	16.22	12.62	*	-
PLACE1010944	83.2	92.75	68.43	51.82	78.83	47.87	87	58.47	73.16	*	-
PLACE1010947	33.76	22.26	22.3	12.84	13.89	8.22	22.66	11.99	13.89	*	-
PLACE1010954	10.33	11.3	11.05	5.3	5.81	5.51	5.34	5.12	4.22	**	**
PLACE1010960	15.62	30.44	20.31	7.94	8.28	7.04	11.37	9.05	8.6	*	-
PLACE1010965	13.17	14.34	9.79	7.54	4.78	3.78	4.15	4.73	5.18	*	-
PLACE1010988	18.41	14.95	14.27	6.84	9.6	5.59	12.82	6.34	9.37	**	*
PLACE1010978	57.65	54.74	38.05	40.68	34.09	13.7	18.85	10.36	20.32	*	-
PLACE1010982	21.44	28.06	15.9	7.42	8.54	11.09	10.56	7.85	9.32	*	-
PLACE1010990	327.93	344.55	256.16	258.07	387.78	240.64	208.45	171.08	173.44	*	-
PLACE1011017	109.18	59.4	54.89	23.72	27.4	15.8	81.94	75.45	99.74	*	-
PLACE1011019	13.37	8.64	8.48	8.37	5.37	4.79	6.61	7.41	7.81	*	-
PLACE1011028	17.87	16.01	13.69	8.74	11.48	8.93	15.71	13.37	15.94	*	-
PLACE1011032	5.5	8.88	7.51	2.84	5.75	2.99	4.5	3.38	3.21	*	-
PLACE1011041	11.32	11.72	11.41	4.79	6.17	4.9	2.74	0	1.85	**	**
PLACE1011045	30.68	27.07	24.09	20.73	22.77	14.24	25.88	22.95	30.08	*	-
PLACE1011046	10.75	10.03	9.15	3.68	6.42	4.38	4.21	3.5	2.56	**	**
PLACE1011054	23.11	28.67	18.54	13.24	17.45	11.68	27.3	21.77	27.93	*	-
PLACE1011056	5.67	3.31	3.16	3.73	4.11	8.48	6.33	3.05	6.91	*	-
PLACE1011057	10.04	10.53	7.99	8.15	6.14	3.72	11.95	6.55	7.6	*	-
PLACE1011059	19.4	17.98	11.33	7.04	9.45	9.52	5.53	6.86	9.19	*	-
PLACE1011066	9.28	7.31	5.97	5.75	7.03	4.1	2.13	5.18	0.49	*	-
PLACE1011087	26.84	40.15	33.04	-67.32	27.48	18.29	127.9	19.42	37.4	*	-
PLACE1011090	10.52	7.52	8.48	3.01	5.18	3.1	5.04	4.61	6.16	*	-
PLACE1011109	43.09	16.58	13.76	6.5	19.94	4.64	12.2	7.21	13.97	*	-
PLACE1011114	17.98	24.04	19.89	13.12	19.15	8.13	12.09	13.25	14.96	*	-
PLACE1011116	30.03	28.53	37.37	10.08	8.87	7	8.26	10.18	5.99	**	**
PLACE1011122	5.9	9.6	5.93	4.16	4.82	3.11	3.97	4.69	2.89	*	-
PLACE1011133	6.47	4.35	6.51	3.81	3.93	3.81	2.72	5.84	2.91	*	-
PLACE1011134	5.48	5.99	6.34	4.28	4.27	3.64	3.72	3.88	3.78	**	**
PLACE1011143	4.78	9.87	5.59	2.21	5.45	3.27	2.93	3.63	4.13	*	-
PLACE1011146	10.48	13.58	12.48	9.3	11.11	4.11	4.38	6.69	4.48	**	-
PLACE1011180	7.71	9.89	6.71	4.98	7.77	3.25	6.45	4.47	3.78	*	-
PLACE1011185	21.99	27.95	32.58	8.87	19.75	11.93	12.6	15.59	9.42	*	-
PLACE1011181	5.25	3.37	6	2.06	4.49	2.29	2.81	4.09	3.85	*	-
PLACE1011186	60.3	38.78	38.17	37.68	34.15	24.31	17.32	33.87	29.06	*	-
PLACE1011188	25.07	30.45	27.42	12.69	29.01	15.83	19.99	27.9	21.4	*	-
PLACE1011203	10.27	18.2	13.8	9.36	12.5	8.63	8.02	13.49	10.02	*	-
PLACE1011214	7.39	9.12	10.92	5.57	5.78	3.85	1.49	2.02	3.84	*	-
PLACE1011219	7.03	5.85	7.27	8.39	4.52	4.58	9.07	4.64	3.23	*	-
PLACE1011221	10.63	11.42	9.42	5.44	10.51	4.82	4.92	5.52	5.28	**	**
PLACE1011229	13.78	13.72	12.78	9.23	7.48	4.66	5.43	7.1	4.99	**	**
PLACE1011231	34.58	28.7	50.92	23.03	24.8	13.83	19.12	24.97	28.97	*	-
PLACE1011238	13.02	10.94	13.95	5.98	7.59	5.06	4.42	6.98	4.82	**	**
PLACE1011247	571.8	409.73	639.71	528.63	738.01	568.23	553.23	542.58	267.58	*	-
PLACE1011283	4.49	7.28	9.65	4.09	5.29	4.39	2.77	4.83	3.37	*	-
PLACE1011273	9.77	15.11	15.29	8	7.91	4.87	6.04	5.28	3.97	*	-
PLACE1011278	21.08	26.71	20.68	8.67	19.17	11.18	9.31	5.1	7.74	**	-
PLACE1011289	8.76	9.56	9.49	5.47	7.47	4.38	4.05	4.14	4.61	*	-
PLACE1011291	3.27	1.73	2.37	1.38	1.37	0.89	2.52	4.22	1.75	*	-
PLACE1011296	15.1	14.79	20.48	7.88	6.09	7.28	12.72	18.07	15.96	**	-
PLACE1011310	49.57	40.91	47.27	26.08	24.65	29.49	26.25	31.46	34.3	**	-
PLACE1011311	12.85	11.18	13.64	5.84	8.4	7.88	4.02	7.29	3.89	**	**
PLACE1011321	318.11	225.18	251.29	230.23	241.81	208.08	340.56	284.43	231.07	*	-

2717

2718

PLACE1011325	67.40	68.67	34.36	32.79	55.65	25.52	26.06	16.66	23.25	*	-
PLACE1011332	37.6	32.99	20.69	24.89	22.84	17.08	26.8	25.99	39.38	*	-
PLACE1011340	6.88	4.22	8.9	7.42	3.33	10.93	4.69	3.22	3.76	*	-
PLACE1011353	13.06	9.45	9.31	4.05	5.07	3.21	9.83	8.83	10.17	**	-
PLACE1011360	7.09	6.29	7.09	5.01	4.64	2.08	4.25	9.92	6.13	*	-
PLACE1011364	36.2	43	48.69	18.82	32.17	16.40	29.76	23.08	23.29	*	-
PLACE1011365	56.47	40.08	43.45	30.4	25.78	22.59	35.29	38.44	44.51	*	-
PLACE1011371	8.03	11.85	15.3	3.37	4.12	3.15	3.67	7.88	4.18	*	-
PLACE1011375	7.04	6.27	10.93	2.06	3.1	2.3	1.72	3.77	5.21	*	-
PLACE1011385	164.63	196.07	182.53	143.65	164.65	123.82	185.62	179.87	160.52	*	-
PLACE1011399	13.53	14.93	12.77	5.31	8.23	5.03	14.48	12.63	13.08	**	-
PLACE1011405	24.35	18.38	25.14	17.24	17.93	13.01	14.05	18.63	13.24	*	-
PLACE1011407	11.98	8.15	11.8	4.39	4.52	4.88	4.7	7.06	4.99	**	-
PLACE1011419	45.6	33.74	47.01	28.72	27.65	25.25	33.12	24.14	29.66	*	-
PLACE1011433	12.4	6.05	13.11	5.89	9.23	4.52	4.03	6.39	4.28	*	-
PLACE1011440	22.91	21.43	16.68	17.18	8.65	6.34	4.22	2.79	3.13	**	-
PLACE1011452	21.99	27.18	21.9	7.47	6.58	8.33	1.98	2.65	3.76	**	-
PLACE1011465	8.69	10.4	8.63	4.43	3.74	3.61	1.27	3.42	2.9	**	-
PLACE1011472	12.35	11.03	8.39	5.78	6.69	4.74	4.12	4.06	4.53	*	-
PLACE1011477	6.89	4.15	8.42	3.82	3.1	2.49	5.68	5.35	7.93	*	-
PLACE1011478	147.6	127.45	151.17	88.45	129.14	92.81	97.94	88.28	67.65	**	-
PLACE1011482	6.88	4.84	8.71	5.69	3.6	3.44	4.78	2.62	3.91	*	-
PLACE1011488	11.88	13.27	11.87	6.42	7.11	5.44	7.49	6.76	6.34	**	-
PLACE1011501	5.51	5	4.78	1.97	2.88	1.92	1.63	1.09	0.63	**	-
PLACE1011503	8.38	7.18	8.72	3.18	3.42	2.41	3.54	1.95	1.63	**	-
PLACE1011509	6.79	8.76	6.6	2.67	3.22	3.57	3.54	1.66	2.48	**	-
PLACE1011514	39.39	42.6	35.59	29.05	24.04	21.59	28.05	21.82	28.44	**	-
PLACE1011516	2.25	2.2	3.02	1.65	1.39	0.92	2.22	1.19	1.28	*	-
PLACE1011520	56.28	54.22	44.97	60.14	59.13	48.68	37.37	33.63	31.08	*	-
PLACE1011538	3.92	2.67	3.89	1.41	2.25	0.89	1.63	0.62	1.52	*	-
PLACE1011555	6.04	5.36	4.18	1.71	1.41	2.12	2.42	1.58	2.13	**	-
PLACE1011561	40.83	32.97	31.05	28.42	20.97	20.97	15.08	18.08	18.45	**	-
PLACE1011563	10.69	8.43	8.69	4.39	3.31	1.28	2.74	2.46	2.05	*	-
PLACE1011567	8.99	5.72	8.65	4.85	2.48	0.84	3.39	1.71	1.08	*	-
PLACE1011569	17.38	15.57	12.39	8.98	14.17	5.41	4.61	3.24	6.59	**	-
PLACE1011578	8.4	4.88	4.15	2.79	3.81	3.52	4.89	6.65	7.51	*	-
PLACE1011585	11.28	10.02	9.62	6.88	3.58	6.49	4.18	5.75	5.57	**	-
PLACE1011595	30.83	17.84	23.55	13.98	14.55	6.78	13.1	10.08	12.82	*	-
PLACE1011641	17.21	13.31	15.6	11.83	10.3	8.13	12.16	8.31	8.02	*	-
PLACE1011642	14.04	9.07	8.81	4.73	3.8	2.88	2.08	3.09	1.68	*	-
PLACE1011643	21.88	15.91	14.82	8.69	7.73	4.5	12.37	9.82	7.66	*	-
PLACE1011645	50.88	55.98	42.82	28.54	24.58	20.05	12.83	13.88	9.31	**	-
PLACE1011649	101.51	209.33	140.08	111.98	127.38	61.19	33.85	36.5	31.67	*	-
PLACE1011650	21.23	18.3	17.27	8.34	7.58	6.35	13.14	8.77	13.14	**	-
PLACE1011661	38.64	29.8	21.88	17.54	11.85	18.02	29.9	27.07	30.09	*	-
PLACE1011664	4693.7	3704.8	3103.3	1925.9	2495.7	2880.1	3091.7	3779.5	2474.5	*	-
PLACE1011672	14.58	19.19	17.65	4.82	11.54	6.52	8.71	8.37	8.68	*	-
PLACE1011675	27.27	28.5	23.64	11.76	7.61	13.02	10.47	6.83	11.83	**	-
PLACE1011682	31.95	30.83	21.79	24.47	12.69	9.18	19.26	16.69	17.58	*	-
PLACE1011708	240.54	265.47	223.19	188.79	260.35	189.88	147.11	130.97	144.28	**	-
PLACE1011719	2115.1	1324.5	1826.8	1953.8	1265.9	1000.2	1428.6	860.26	1289.9	*	-
PLACE1011725	14.63	11.44	8.67	4.74	8.3	2.04	9.22	9.94	8.53	*	-
PLACE1011729	17.92	12.45	10.82	5.48	6.85	3.57	9.59	6.92	10.12	*	-
PLACE1011741	19.11	18.45	8.97	6.48	6.39	4.13	5.03	4.62	6.01	*	-
PLACE1011749	188.63	245.38	188.1	88.37	138.73	83.5	148.84	106.95	169.58	*	-
PLACE1011757	55.97	64.31	51.13	22.6	27.43	20.99	59.9	47.98	65.48	**	-
PLACE1011762	112.55	118.61	84.99	88.65	83.7	57.21	113.37	92.71	109.2	*	-
PLACE1011778	14.17	14.57	14.58	9.09	8.23	7.54	9.58	8.34	17.29	**	-
PLACE1011783	34.3	38.67	23.05	14.78	12.18	17.09	15.47	21.82	20.45	*	-
PLACE1011795	28.68	17.88	14.49	8.41	7.89	4.41	13.18	9.37	14.58	*	-
PLACE1011810	8.48	2.52	3.35	2.53	2.07	0.8	3.19	3.08	3.38	*	-
PLACE1011824	88.62	55.93	50.79	29.92	21.66	12.01	21.62	24.58	19.64	*	-
PLACE1011825	47.11	53.31	45.03	28.52	34.47	24.16	61.42	44.87	47	**	-
PLACE1011835	12.62	11.42	11.32	6.55	7.84	4.84	7.39	8.98	8.87	**	-

140 / 157

2719

2720

PLACE1011836	18.92	14.13	10.67	5.56	9.72	5.44	13.34	8.9	14.01	*	-	-
PLACE1011847	18.39	16.08	12.62	9.39	11.06	5.84	18.15	10.26	11.1	*	-	-
PLACE1011856	44.57	41.99	25.3	17.11	33.33	14.76	55.6	42.68	40.67	*	-	-
PLACE1011868	17.26	16.38	17.11	5.33	7.08	6.53	8.22	8.74	6.1	**	**	-
PLACE1011874	23.12	12.72	25.71	18.84	19.22	7.85	10.17	8.71	12.51	*	-	-
PLACE1011876	10.58	9.98	11.17	5.18	10.63	6.87	7.18	8.04	4.93	**	-	-
PLACE1011877	4.39	5.14	4.95	3.7	5.62	3.78	2.69	4.05	4.24	*	-	-
PLACE1011891	21.71	17.57	18.98	7.39	14.47	8.67	8.21	9.15	7.88	*	-	-
PLACE1011896	15.35	16.64	24.3	10.41	10.14	12.14	5.01	8.13	7.98	*	-	-
PLACE1011929	16.12	14.42	15.8	7.68	8.85	7.76	8.48	2.46	3.24	**	**	-
PLACE1011922	9.14	7.55	8.03	4.49	8.39	4.08	7.97	6.15	5.33	*	-	-
PLACE1011923	166.88	139.43	148.9	104.2	117.42	86.95	136.93	140	124.61	*	-	-
PLACE1011937	11.45	9.18	10.8	8.58	6.75	5.1	5.71	6.29	5.45	*	-	-
PLACE1011939	20.33	21.55	26.93	12.79	15.19	13.44	14.68	11.31	12.67	*	-	-
PLACE1011940	46.38	48.76	46.73	35.11	34.53	31.67	43.31	41.44	39.86	**	**	-
PLACE1011962	58.59	70.73	80.18	30.15	38.09	28.84	47.96	47.88	42.57	**	-	-
PLACE1011984	53.78	49.36	48.95	41.43	47.33	38.34	22.23	20.62	20.74	**	-	-
PLACE1011978	50.21	52.88	64.71	34.8	31.39	22.36	22.48	20.19	18.12	*	-	-
PLACE1011980	13.57	18.54	10.96	9.65	15.28	6.63	4.88	9.99	8.31	*	-	-
PLACE1011981	45.23	28.89	45.44	16.88	15.91	9.80	14.66	11.01	11.18	*	-	-
PLACE1011982	116.69	99.4	123.68	57.42	68.6	45.81	97.45	98.55	115.07	**	-	-
PLACE1011995	8.31	11.27	12.33	6	7.29	7.22	8.27	8.29	4.87	*	-	-
PLACE1012026	36.38	36.13	39.18	21.48	26.03	23.3	20.5	21.06	24.79	**	**	-
PLACE1012026	72.98	80.46	66.75	52.71	40.29	50.42	21.72	33.87	25.71	**	**	-
PLACE1012031	8.69	7.34	9.88	2.73	6.11	3.89	3.41	3.11	3.97	*	-	-
PLACE2000003	144.76	222.75	62.64	74.47	133.88	40.79	39.01	32.21	30.64	*	-	-
PLACE2000005	20.72	25.78	18.16	9.79	22.43	11.52	5.78	6.65	14.3	*	-	-
PLACE2000006	74.92	61.78	61.3	27.4	24.75	21.12	60.02	43.94	72.77	**	-	-
PLACE2000007	11.31	11.29	12.53	6.68	7.02	5.2	5.63	10.02	7	**	-	-
PLACE2000011	20.07	19.35	21.84	14.92	4.68	16.77	9.39	11.52	19.43	*	-	-
PLACE2000014	10.47	6.72	12.3	4.76	6.02	3.48	3.63	8.9	4.73	*	-	-
PLACE2000015	63.77	83.22	43.82	34.08	43.14	43.39	19.29	26.64	38.8	*	-	-
PLACE2000017	64.44	66.07	78.07	40.68	48.09	29.95	59.7	61.53	44.91	*	-	-
PLACE2000021	131	135.36	142.32	116.28	142.66	101.82	89.75	68.71	88.35	**	-	-
PLACE2000022	6.62	6.76	7.62	3.85	5.14	3.55	4.57	1.84	5.72	**	-	-
PLACE2000030	277.16	275.48	252.39	182.81	167.16	156.31	125.23	96.07	135.66	**	-	-
PLACE2000032	8.03	6.4	8.8	4.38	5.33	2.81	4.32	15.18	4.01	*	-	-
PLACE2000033	19.24	16.61	20.22	9.43	14.03	10.83	8.93	12.91	10.87	*	-	-
PLACE2000034	212.75	182.06	146.24	218.9	195.88	170.13	146.86	112.34	126.26	*	-	-
PLACE2000039	8.9	10.84	13.51	4.94	3.89	4.8	3.72	4.38	6.52	**	-	-
PLACE2000043	13.64	20.44	15.11	6.28	13.12	6.04	6.05	7.64	6.22	*	-	-
PLACE2000044	7.28	6.48	9.44	3.48	5.71	4.79	4.48	4.25	4.59	*	-	-
PLACE2000047	17.41	19.39	12.83	5.83	12.83	7.05	7.81	8.54	5.78	*	-	-
PLACE2000050	18.34	17.87	22.05	7.34	8.88	7.31	7.51	8.31	7.94	**	-	-
PLACE2000081	59.75	61.97	78.23	34.92	41.03	37.88	22.68	14.04	20.77	*	-	-
PLACE2000082	44.19	44.8	62.08	41.54	45.17	33.58	38.88	38.08	40.24	*	-	-
PLACE2000072	218.69	181.79	181.98	280.73	185.27	186.23	111.76	113.44	105.63	**	-	-
PLACE2000073	8.16	10.75	9.27	3.77	4.89	3.62	4.58	4.79	3.59	**	-	-
PLACE2000097	128.94	128.07	120.9	148.48	131.74	105.73	80.14	87.25	86.70	**	-	-
PLACE2000100	19.48	14.19	15.99	6.02	8.77	5.56	4.8	3.51	6.24	**	-	-
PLACE2000103	365.88	328.95	197.42	251.22	438.52	235.01	231.23	338.03	237.95	*	-	-
PLACE2000106	32.23	23.41	23.14	14.48	8.17	13.03	9.13	10.5	5.27	*	-	-
PLACE2000111	162.37	157.38	125.3	88.8	105.91	83.66	90.52	44.23	69.19	*	-	-
PLACE2000115	27.01	16.97	16.32	19.6	21.67	16.26	13.74	11.32	11.63	*	-	-
PLACE2000118	229.23	214.41	142.27	124.91	162.19	92.16	153.7	62.85	67.07	*	-	-
PLACE2000124	7.33	5.71	5.57	2.12	1.78	1.65	2.01	1.58	1.09	**	-	-
PLACE2000132	18.07	19.01	19.11	15.64	11.18	12.49	9.7	6.52	8.48	*	-	-
PLACE2000136	33.6	40.68	26.33	13.95	42.68	22.64	9.98	17.8	18.28	*	-	-
PLACE2000137	6.87	5.34	6.91	1.81	2.95	2.09	1.24	1.8	2.36	**	-	-
PLACE2000140	11.37	11.51	12.2	8.07	9.45	9.49	11.48	11.25	10.8	**	-	-
PLACE2000147	10.22	7.24	9.46	6.01	5.85	4.51	7.52	7.8	9.27	*	-	-
PLACE2000153	17.95	21.42	13.85	5.61	8.67	8.43	4.84	5.27	2.27	*	-	-
PLACE2000164	8.53	12.11	9.44	3.18	4.26	4.12	4.98	3.97	2.13	**	-	-
PLACE2000170	38.84	46.93	33	28	28.66	22.27	30.03	23.36	30.82	*	-	-

141 / 157

2721

2722

PLACE2000172	9.55	12.02	10.88	4.88	5.87	3.32	4.01	4.8	8.2	00	00	-	-
PLACE2000173	16.63	20.86	19.04	9.66	11.13	8.7	6.74	8.23	3.14	00	00	-	-
PLACE2000174	8.73	7.76	6.4	3.37	4.73	2.4	3.08	3.28	1.62	0	00	-	-
PLACE2000176	47.07	39.82	36.01	26.8	32.78	24.97	24.68	26.79	25.41	0	00	-	-
PLACE2000187	11.49	9.44	7.22	4.28	4.58	4.45	2.47	4.64	3.84	0	0	-	-
PLACE2000218	16.24	13.28	18.1	7.83	18.53	7.67	11.12	11.23	10.67	00	0	-	-
PLACE2000219	49.32	28.21	37.3	18.17	8.67	13.04	7.68	7.79	8.4	0	00	-	-
PLACE2000221	9.58	7.64	7.48	2.22	3.99	1.24	4.08	2.43	3.07	00	00	-	-
PLACE2000223	16.67	12.34	12.85	4.95	8.75	4.32	4.43	2.01	2.87	00	00	-	-
PLACE2000231	8.48	8.65	7.01	3.78	6.92	4.27	4.19	2.32	3.59	0	00	-	-
PLACE2000236	7.02	5.03	8.86	1.64	5.33	1.9	4.43	3.57	3.78	0	00	-	-
PLACE2000246	21.31	17.8	13.21	7.9	13.4	6.47	11.52	8.8	10.91	0	0	-	-
PLACE2000284	41.11	43.3	30.68	20.28	22.27	18.14	15.54	15.85	15.69	0	00	-	-
PLACE2000274	6.8	7.42	7.38	4.35	3.04	1.89	3.84	3.12	4.31	00	00	-	-
PLACE2000287	32.11	30.76	24.89	20.39	26.81	21.53	17.98	16.3	19.45	00	0	-	-
PLACE2000298	42.94	34.68	35.83	23.36	22.17	20.17	32.02	14.36	29.35	00	0	-	-
PLACE2000302	252.74	304.77	237.45	285.35	287.99	217.29	124.09	112.58	108.48	00	00	-	-
PLACE2000305	21.47	23.58	17.69	18.14	17.23	16.65	22.14	18.21	22.01	0	0	-	-
PLACE2000317	37.2	38.12	23.24	20.01	28.38	14.37	31.05	33.18	31.64	0	0	-	-
PLACE2000324	18.44	17.84	10.35	6.13	8.75	4.88	3.24	8.74	3.2	0	0	-	-
PLACE2000334	21.38	17.8	11.43	10.36	9.7	9.37	7.54	6.15	8.1	0	00	-	-
PLACE2000335	71.59	65.59	64.82	51.15	35.99	40.68	33.77	38.89	29.67	0	00	-	-
PLACE2000340	20.82	28.95	19	17.88	23.88	12.85	16.23	14.73	18.82	00	0	-	-
PLACE2000341	14.8	11.92	10.45	6.64	5.34	5.33	5.38	5.46	7.68	00	0	-	-
PLACE2000342	23.63	21.95	21.87	16.28	11.88	11.51	22.81	14.69	23.12	00	00	-	-
PLACE2000347	9.67	9.29	10.05	4.16	2.94	3.3	5.78	3.54	5.48	00	00	-	-
PLACE2000357	112.17	70.59	69.73	83.45	100.62	78.25	23	68.79	48.64	00	00	-	-
PLACE2000368	67.37	32.59	34.74	25.22	20.69	15.48	44.83	47.18	42.01	00	00	-	-
PLACE2000369	86.39	48.68	57.2	40.91	54.54	32.45	1.96	27.1	55	00	00	-	-
PLACE2000386	24.09	23.36	18.68	12.65	12.71	9.05	11.76	8.72	10.83	00	00	-	-
PLACE2000371	12.17	9.37	8.13	5.98	7.78	4.49	7.64	10.97	8.2	00	00	-	-
PLACE2000373	8.3	7.58	8.83	4.52	8.25	4.81	3.89	1.17	2.44	00	00	-	-
PLACE2000374	20.82	21.67	18.21	18.3	16.71	14.02	11.71	11.44	14.19	0	00	-	-
PLACE2000379	189.97	110.55	125.02	98.68	141.6	89.14	84.84	85.75	71.64	00	00	-	-
PLACE2000386	27.43	27.19	20.99	10.83	12.4	11.22	15.3	17.99	10.26	00	0	-	-
PLACE2000388	260.97	170.84	203.4	172.65	236.63	173.11	130.31	109.84	93.57	0	0	-	-
PLACE2000392	241.74	206.31	271.34	159.33	189.87	145.83	119.67	115.9	95.8	0	00	-	-
PLACE2000394	10.8	10.42	13.61	8.59	15.25	7.68	5.69	8.12	6.51	0	0	-	-
PLACE2000398	16.83	19.82	17.04	9.21	12.98	10.99	7.6	16.14	8.57	00	0	-	-
PLACE2000399	30.8	27.17	17.03	25.83	27.41	28.06	9.08	19.74	20.17	00	00	-	-
PLACE2000402	5.25	8.04	8.31	4.2	8.81	3.39	4.84	3.89	6.41	00	00	-	-
PLACE2000404	24.79	49.52	29.14	49.74	21.64	26.22	16.23	18.08	8.51	00	00	-	-
PLACE2000411	5.67	8.96	8.76	7.21	11.84	6.13	5.19	4.54	7.08	00	00	-	-
PLACE2000418	10.3	8.94	9.87	6.1	3.09	4.67	4.68	5.3	4.11	00	00	-	-
PLACE2000419	98.71	94.48	148.23	68.48	67.97	27.06	44.37	48.85	45.47	0	0	-	-
PLACE2000425	9.83	11.3	13.49	7.49	11.28	8.24	5.84	7.49	5.62	0	0	-	-
PLACE2000427	16.21	22.92	21.67	15.7	21.58	15.92	5.81	6.9	4.78	00	00	-	-
PLACE2000433	26.58	27.32	28.12	13.97	18.4	14.15	21.62	27.64	18.08	00	00	-	-
PLACE2000435	16.36	18.11	21.19	9.69	13.78	7.6	8.43	10.79	12.73	0	0	-	-
PLACE2000438	35.17	38.25	36.72	24.14	34.38	15.98	38.19	37.23	26.32	00	00	-	-
PLACE2000450	24.38	41.59	23.11	16.35	37.33	18.21	23.47	16.65	30.25	0	00	-	-
PLACE2000455	51.05	47.04	53.8	30.31	38.57	21.71	39.12	34.02	35.42	0	00	-	-
PLACE2000458	13.07	11.68	13.21	8.82	7.75	7.03	6.8	8.78	5.76	00	00	-	-
PLACE2000484	9.95	10.55	11.94	7.59	6.37	5.51	7.66	8.47	7.18	00	0	-	-
PLACE2000485	182.2	189.36	208.18	168.02	148.04	158.91	139.29	234.97	151.61	0	0	-	-
PLACE2000473	23.65	24.51	24.43	18.65	18.15	16.44	24.18	30.07	29.1	00	00	-	-
PLACE2000477	14.25	12.47	18.65	9.89	12.47	8.23	10.6	9.09	8.9	0	0	-	-
PLACE2000084	49.49	41.47	56.9	42.83	53.35	39.93	39.48	28.43	32.45	0	0	-	-
PLACE2000089	30.13	36.74	28.18	19.79	35.05	20.31	13.64	15.44	20.20	00	00	-	-
PLACE2000020	27.99	24.44	23.74	13.45	15.64	8.65	14.29	14.44	13.63	00	00	-	-
PLACE2000029	13.95	12.7	15.35	6.3	9.27	7.3	4.68	16.59	5.52	00	00	-	-
PLACE2000036	8.52	4.32	7.89	5.63	3.97	4.12	2.88	6.95	6.18	00	00	-	-
PLACE2000052	8.21	5.21	10.48	3.53	4.29	4.15	2.75	5.07	2.29	0	0	-	-
PLACE2000059	7.39	8.1	10.8	4	5.08	2.55	2.83	4.28	4.31	0	0	-	-

142 / 157

2723

2724

PLACE300087	49.68	45.52	57.37	38.78	41.13	26.18	39.8	48.08	41.69				
PLACE300088	22.6	20.15	23	9.32	14.75	10.61	1.9	9.3	14.18	**	*	-	-
PLACE300089	28.38	29.24	22.17	14.93	25.34	18.73	8.82	18.65	18.21	*	*	-	-
PLACE300090	7.35	7.9	9.44	5.25	4.6	6.13	2.42	4.32	5.29	*	*	-	-
PLACE300091	35.89	28.1	30.76	15.16	14.79	15.09	26.05	30.71	30.91	**	*	-	-
PLACE300092	42.11	31.82	37.82	38.13	32.18	27.02	30.65	23.15	27.65	*	*	-	-
PLACE300093	419.8	389.05	314.65	379.74	412.05	282.08	411.58	374.58	344.25				
PLACE300094	20.62	25.87	28.77	15.03	9.64	8.12	18.24	11.89	14.6	**	*	-	-
PLACE300095	7.68	8.11	13.16	2.72	3.39	4.18	4.62	3.19	1.88	*	*	-	-
PLACE300096	24.15	28.81	28.5	11.04	24.59	15.08	9.37	8.94	11.68	**	*	-	-
PLACE300097	18.45	12.13	16.32	7.23	18.62	6.93	8.46	9.12	6.78	*	*	-	-
PLACE300098	109.42	82.46	138.9	40.47	87.42	32.89	57.71	63.5	68.14				
PLACE300099	68.44	59.97	67.42	62.74	55.88	38.34	37.83	26.14	38.79	**	*	-	-
PLACE300100	10.3	11.14	12.88	7.95	8.64	6.56	7.31	6.3	7.57	*	**	-	-
PLACE300101	54.28	50.77	22.77	23.49	28.34	24.7	10.12	11.93	8.34	*	*	-	-
PLACE300102	9.77	8.45	8.44	2.87	5.38	4.07	2.62	2.18	2.82	*	**	-	-
PLACE300103	38.44	50.52	35.49	23.64	23.38	25.04	21.36	24.91	20.01	*	*	-	-
PLACE300104	10.22	9.34	10.13	4.45	4.83	3.36	2.93	2.26	3.02	**	**	-	-
PLACE300105	19.92	29.65	15.83	8.71	11.7	10.32	5.77	7.29	4.67	*	*	-	-
PLACE300106	4.65	6.88	7.85	4.88	4.67	2.21	2.88	2.72	3.49	*	*	-	-
PLACE300107	12.63	9.87	13.48	6.15	8.69	4.33	5.58	7.04	5.47	**	**	-	-
PLACE300108	1627	2636.2	1913.4	1710.4	2608.8	2399	3320.1	2828.5	2357.1				
PLACE300109	7.18	4.55	8.38	2.14	1.71	1.68	2.79	2.24	2.67	*	*	-	-
PLACE300110	19.48	15.17	17.17	4.88	8.73	7.24	5.33	5.58	6.07	**	**	-	-
PLACE300111	12.53	18.93	16.84	6.91	6.83	4.67	7.28	5.87	4.15	**	**	-	-
PLACE300112	7.97	8.69	7.83	2.48	4.45	2.85	2.71	2.19	2.08	**	**	-	-
PLACE300113	5.49	5.6	6.62	1.88	8.49	2.8	2.64	3.52	2.48	**	*	-	-
PLACE300114	40.02	27.48	31.62	27.29	30.91	23.89	26.59	30.58	23.2				
PLACE300115	20.76	15.1	17.45	10.19	6.64	8.25	15.48	14.18	17.02	**	*	-	-
PLACE300116	13.39	10.52	10.44	6.03	3.81	4.29	4.65	4.57	4.68	**	**	-	-
PLACE300117	61.41	58.02	42.84	38.7	37	28.53	36.15	41.5	44.54	*	*	-	-
PLACE300118	12.37	9.42	8.64	5.7	3.63	2.33	5.09	3.54	2.87	*	**	-	-
PLACE300119	180.88	144.07	131.68	178.14	175.28	127.3	77.77	64.65	88.2	**	**	-	-
PLACE300120	20.28	21.69	18.74	7.11	11.09	8.3	3.48	2.23	3.79	**	**	-	-
PLACE300121	17.48	18.88	14.84	8.27	14.11	10.52	4.85	3.25	5.88	**	*	-	-
PLACE300122	17.8	13.68	8.85	9.08	2.29	4.85	6.33	8.35	7.81				
PLACE300123	9.81	7.95	8.28	4.48	5.85	4.02	5.72	2.93	4.77	**	*	-	-
PLACE300124	58.11	27.4	48.85	17.82	13.78	17.4	11.61	10.29	11.25	*	*	-	-
PLACE300125	18.63	13.18	20	8.6	7.54	7.64	10.16	10.98	16.61	*	*	-	-
PLACE300126	68.99	47.97	42.85	32.37	25.93	24.8	32.88	24.78	32.96	*	*	-	-
PLACE300127	27.82	25.4	19.81	15.47	18.42	14.64	15.57	11.42	13.74	*	*	-	-
PLACE300128	14.58	20.8	13.28	5.71	8.52	5.71	8.32	6.43	8.38	*	*	-	-
PLACE300129	13.09	13.68	8.83	2.78	7.9	3.42	3.8	5.28	3.1	*	**	-	-
PLACE300130	37.31	49.14	39.5	28.72	17.9	18.28	33.17	28.5	40.23	*	*	-	-
PLACE300131	12.29	9.64	9.42	6.2	5.6	4.27	5.42	4.79	6.47	**	**	-	-
PLACE300132	51.34	49.08	45.31	11.5	35.47	14.78	19.68	8.79	16.35	*	**	-	-
PLACE300133	24.3	28.61	23.33	14.31	29.51	12.27	15.7	15.03	15.45	*	**	-	-
PLACE300134	43.24	35.12	33.82	31.65	31.68	28.75	27.52	20.55	23.58	*	*	-	-
PLACE300135	28.37	27.71	24.43	12.33	16.5	11.35	20.86	17.37	21.33	**	*	-	-
PLACE300136	6.68	8.07	7.87	8.55	6.13	3.84	2.57	2.99	0.04	**	**	-	-
PLACE300137	393.24	473.08	270.29	359.87	515.03	306.41	470.48	682.39	717.67				
PLACE300138	11.78	5.98	6.82	3.79	5.41	2.48	9.74	10.29	11.53				
PLACE300139	12.2	6.88	6.99	4.2	5.65	4.02	4.35	5.75	4.94				
PLACE300140	13.71	12.65	10.21	4.86	8.13	4.94	5.77	4.37	7.49	*	*	-	-
PLACE300141	17.23	29.11	20.8	8.42	15.27	8.16	11.43	7.14	13.58	*	*	-	-
PLACE300142	14.74	14.86	16.82	8.78	8.26	6.88	10.37	5.48	6.79	**	**	-	-
PLACE300143	57.42	93.8	72.53	67.02	58.59	35.38	59.88	38.27	60.8				
PLACE300144	9.44	8.81	10.65	4.39	5.19	4.3	7.08	4.68	8.58	**	*	-	-
PLACE300145	6.59	7.45	4.94	1.48	2.65	3.2	4.78	2.37	2.95	*	*	-	-
PLACE300146	38.45	30.27	25.67	4.05	7.88	3.83	14.18	11.37	15.74	**	*	-	-
PLACE300147	48.65	44.25	34.93	16.2	28.01	11.78	24.23	27.25	23.51	*	*	-	-
PLACE300148	5.55	3.77	3.38	1.68	3.57	0.43	3.58	2.95	3.21	*	*	-	-
PLACE300149	28.03	22.21	15.9	9.88	12.79	6.37	21.94	28.7	24.22	*	*	-	-
PLACE300150	15.37	13.07	9.83	5.64	4.9	3.93	10.68	11.05	9.52	**	*	-	-

143/157

2725

2726

PLACE3000400	42.78	33.62	30.29	11.43	28.51	8.47	52.78	33.47	38.48	*	-	-
PLACE3000401	137.51	137.03	85.08	81.28	111.28	51.38	61.68	72.2	62.85	*	-	-
PLACE3000402	8.43	7.72	4.24	2.44	6.58	2.29	8.31	7.96	5.44	*	-	-
PLACE3000403	7.33	4.67	7.44	2.73	3.5	5.1	3.7	6.44	1.68	*	-	-
PLACE3000406	4.67	3.39	6.42	4.46	6.89	4.18	3.34	4.09	2.59	*	-	-
PLACE3000413	12.33	14.38	18.34	10.31	15.31	7.77	6.83	11.34	9.07	*	-	-
PLACE3000416	84.6	58.17	41.71	94.16	128.08	86.62	51.61	89.65	63.01	*	-	-
PLACE3000425	2.87	4.19	4.03	2.67	4.11	2.12	2.08	0.77	1.54	*	-	-
PLACE3000437	8.08	4.62	9.74	3.57	11.5	3	3.57	3.67	4.77	*	-	-
PLACE3000455	6.66	7.45	7.23	5.48	5.38	4.19	5.5	5.74	3.82	*	-	-
PLACE3000475	4.99	6.71	5.08	4.69	6.48	5.91	3.2	5.35	2.87	*	-	-
PLACE3000477	12.61	9.21	10.6	8.81	9.19	10.35	8.13	7.94	5.88	*	-	-
PLACE3000503	30	20.47	39.94	8.43	16.43	9.82	12.11	10.62	8.23	*	-	-
PLACE4000000	8.19	8.19	9.59	5.97	7.85	6.93	4.18	5.67	3.47	**	-	-
PLACE4000009	46.88	36	37.99	37.22	51.39	35.97	11.09	18.85	7.63	**	-	-
PLACE4000014	18.34	18.67	10.39	16.82	18.49	18.69	6.08	13.2	15.87	**	-	-
PLACE4000029	5.3	5.07	5.95	2.91	4.87	2.82	3.88	3.02	2.87	**	-	-
PLACE4000034	10.02	8.01	6.28	5.12	7.93	4.55	5.88	5.27	5.14	**	-	-
PLACE4000049	21.88	18.28	18.64	14.38	19.7	19.05	12.54	13.86	11.06	**	-	-
PLACE4000052	47.83	33.03	40.39	17.98	15.08	20.28	24.01	22.07	24.29	**	-	-
PLACE4000082	53.48	48.6	50.82	32.69	40.48	28.08	39.71	38.99	45	*	-	-
PLACE4000083	58.25	68.62	67.5	55.27	69.37	73.68	45.78	68.53	44.9	*	-	-
PLACE4000089	10.35	13.86	13.41	10.88	10.48	9.25	5.88	6.4	5.88	**	-	-
PLACE4000093	15.33	18.02	19.57	10.49	13.84	10.33	17.17	14.6	14.04	*	-	-
PLACE4000100	8.21	7.32	11.21	7.61	8.2	7.32	5.19	5.2	4.71	*	-	-
PLACE4000101	13.85	15.73	12.84	8.67	14.38	8.43	4.02	8.73	6.65	**	-	-
PLACE4000106	32	18.81	23.23	18.35	21.38	15.88	17.78	20.58	18.06	**	-	-
PLACE4000128	22.06	18.53	23.14	8.88	10.24	9.57	11.18	17.49	11.78	**	-	-
PLACE4000129	268.34	168.01	188.25	120.37	95.41	112.42	146.64	88.31	136.33	*	-	-
PLACE4000131	59.1	45.51	65.67	41.68	34.82	39.78	18.29	19.7	30.83	*	-	-
PLACE4000147	7.89	7.51	11.78	3.51	5.07	3.96	2.77	6.32	2.52	*	-	-
PLACE4000156	15.48	18.75	19.37	9.1	12.85	10.65	15.8	15.48	10.02	*	-	-
PLACE4000175	12.45	10.41	18.08	11.94	9.87	2.48	5.38	8.69	6.89	*	-	-
PLACE4000180	40.84	40.93	32.85	23.06	35.22	28.68	12.42	10.73	11	**	-	-
PLACE4000182	35.69	24.13	23.88	16.77	21.69	17.65	17.75	18.51	18.4	**	-	-
PLACE4000208	33.82	28.03	26.48	11.73	17.07	14.5	17.45	20.18	20.02	**	-	-
PLACE4000211	12.88	13.88	13.86	7.98	9.94	9.19	6.64	31.04	6.54	**	-	-
PLACE4000214	13.12	6.23	9.29	3.48	7.5	4.18	4.98	7.25	13.71	**	-	-
PLACE4000222	35.35	30.73	34.54	24.03	24.44	20.1	20.58	28.37	25.45	**	-	-
PLACE4000223	14.88	14.83	15.03	7.38	18.28	5.52	4.18	7.87	6.85	**	-	-
PLACE4000229	12.38	12.18	12.52	4.51	6.31	4.98	4.32	4.28	3.98	**	-	-
PLACE4000230	9.56	8.77	9.66	4.68	7.08	5.7	4.97	8.82	3.71	**	-	-
PLACE4000233	22.71	24.01	28.85	11.67	13.88	13.77	8.53	11.22	9.26	**	-	-
PLACE4000239	10.07	7.4	9.42	6.8	7.34	6.09	4.49	4.54	3.54	**	-	-
PLACE4000247	15	11.82	15.68	9.41	13.91	8.67	8.67	6.24	7.39	**	-	-
PLACE4000250	18.14	18.27	20.25	16.64	17.88	12.73	10.47	8.04	11.21	**	-	-
PLACE4000252	10.01	5.15	12.75	1.39	3.88	3.1	2.21	5.92	2.7	*	-	-
PLACE4000258	25.72	27.31	18.18	12.78	18.89	8.44	8.29	18.39	9.08	**	-	-
PLACE4000261	23.52	25.16	22.9	11.05	14.2	9.22	13.61	14.62	13.29	**	-	-
PLACE4000264	176.03	166.64	118.78	135.7	191.87	147.27	105.16	95.1	89.72	*	-	-
PLACE4000269	71.18	48.07	82.08	23.03	41.63	24.74	71.71	82.8	52.18	*	-	-
PLACE4000270	7.6	6.37	8.9	4.89	5.92	4.81	3.05	3.68	3.44	*	-	-
PLACE4000281	16.28	16.07	18.29	8.09	10.92	7.61	6.44	6.66	5.1	**	-	-
PLACE4000300	8.43	9.82	8.78	5.3	8.5	7.03	3.68	2.93	2.96	**	-	-
PLACE4000320	12.37	11.22	11.92	6.27	10.09	6.43	4.47	4.26	2.85	*	-	-
PLACE4000323	15.18	16.25	19.18	9.12	14.08	10.9	6.05	4.54	4.03	*	-	-
PLACE4000328	11.15	8.97	10.53	4.35	4.69	4.13	4.28	3.68	4.65	**	-	-
PLACE4000344	27.71	32.93	24.51	14.27	30.44	19.64	5.14	9.78	8.98	**	-	-
PLACE4000347	158.82	129.54	154.28	89.79	129.04	79.83	138.06	137.09	135.75	*	-	-
PLACE4000354	15.47	9.48	11.43	7.01	8.32	6.55	3.28	5.98	4.83	*	-	-
PLACE4000367	8.63	5.47	7.82	5.38	5.39	6.72	3.84	3.95	3.64	*	-	-
PLACE4000369	11.27	6.37	9.31	5.27	4.5	4.05	4.11	4.32	3	*	-	-
PLACE4000379	6.35	6	4.77	3.22	2.93	2.33	3.1	3.32	2.21	**	-	-
PLACE4000387	25.99	34.68	32.08	17.28	20.68	18.67	6.75	8.88	3.97	*	-	-

2727

2728

PLACE4000392	8.31	9.76	11.03	3.63	5.95	3.64	5.73	2.01	2.93	**	-	-
PLACE4000399	53.9	51.54	53.28	20.64	28.55	18.63	37.94	35.1	44.54	**	-	-
PLACE4000401	4.22	3.76	3.44	3.98	3.49	2.6	3.5	3.9	1.61	*	-	-
PLACE4000403	8.42	5.82	6.5	4.19	7.67	4.85	3.98	3.29	3.53	*	-	-
PLACE4000411	20.23	15.5	15.73	8.23	11.19	9.81	14.98	10.1	11.34	*	-	-
PLACE4000415	8.15	4.8	4.61	4.08	6.07	4.23	2.98	3.98	4.5	*	-	-
PLACE4000416	95.43	25.11	24.69	17.72	17.34	16.23	24.55	17.6	25.73	*	-	-
PLACE4000424	43.93	33.38	31.83	13.69	19.37	13.98	27.93	23.64	33	**	-	-
PLACE4000431	10.44	8.78	8.78	5.88	4.85	3.5	4.69	2.33	4.08	**	-	-
PLACE4000443	50.64	44.47	41.43	26.49	37.08	21.33	51.22	33.11	43.74	*	-	-
PLACE4000445	5.42	4.17	3.6	3.57	7.43	5.38	2.23	3.95	2.66	*	-	-
PLACE4000450	9.85	6.68	6.23	4.72	8.84	4.07	3.58	7.58	3.73	*	-	-
PLACE4000455	167.81	132.51	128.81	98.87	124.22	85.79	172.16	139.08	172.24	*	-	-
PLACE4000458	31.84	25.74	18.63	10.79	23.42	11.27	11.28	10.8	17.69	*	-	-
PLACE4000468	13.01	11.54	9.57	6.58	6.92	4.59	5.78	4.65	5.6	*	-	-
PLACE4000472	165.28	158.69	152.44	150.34	164.09	112.33	151.06	118.58	147.01	*	-	-
PLACE4000487	10.38	6.89	9.12	4.68	4.63	2.93	3.52	2.06	3.49	*	-	-
PLACE4000489	8.3	7.97	5.85	3.99	5.8	4.43	4.22	4.11	4.53	*	-	-
PLACE4000494	70.61	55.91	43.58	29.17	31.84	24.84	72.87	73.79	78.08	*	-	-
PLACE4000502	204.58	180.8	176.1	158.92	168.08	118.82	219.32	195.78	200.53	*	-	-
PLACE4000521	28.26	22.58	17.07	7.12	15.76	9	21.36	20.08	19.61	*	-	-
PLACE4000522	18.48	20.38	16.27	10.48	14.51	9.51	18.91	13.88	17.23	*	-	-
PLACE4000537	9.79	11.19	9.86	4.99	8.2	4.43	2.5	3.5	0.58	*	-	-
PLACE4000548	12.35	10.52	9.27	3.9	4.07	4.41	4.08	3.04	3.54	**	-	-
PLACE4000558	56.63	61.59	46.82	23.74	32.06	21.71	65.44	38.24	52.83	**	-	-
PLACE4000581	9.34	8.43	8.7	5.21	3.71	3.17	5.23	4.69	4.75	**	-	-
PLACE4000590	4.28	4.39	2.54	2.29	1.9	1.58	2.69	1.47	1.72	*	-	-
PLACE4000593	9.43	8.29	6.55	6.59	7.86	5.28	9.13	5.16	4.1	*	-	-
PLACE4000612	50.54	44.64	46.86	32.73	32.12	18.75	8.58	0	9.68	*	-	-
PLACE4000638	8.47	6.12	6.11	6.07	5.18	3.4	1.44	4.91	1.37	*	-	-
PLACE4000650	11.71	8	6.58	1.78	5.39	2.69	5.01	3.89	2.3	*	-	-
PLACE4000651	10.02	10.01	7.86	4.63	6.67	3.43	7.21	8.04	6.23	*	-	-
PLACE4000654	40.79	37.09	28.74	21.7	29.39	18.93	39.41	24.87	38.02	*	-	-
PLACE4000670	7.33	7.03	6.77	5.23	9.34	3.97	6.41	3.63	6.87	*	-	-
PLACE4000685	27.36	21.26	28.71	7.25	13.45	14.89	8.25	12.79	8.57	*	-	-
PLACE4000687	5.87	2.72	5.94	3.75	3.52	3.04	2.8	3.66	2.1	*	-	-
PLACE50000003	13.69	15.66	18.88	8.92	15.85	13.17	6.38	8.93	8.84	**	-	-
PLACE50000005	17.05	13.36	13.46	14.47	17.35	13.75	5.61	8.29	7.64	**	-	-
PLACE50000019	9.12	11.38	11.73	6.9	5.85	5.55	3.51	3.39	5.14	**	-	-
PLACE50000021	28.4	29.38	31.86	18.99	18.21	16.52	23.2	21.13	21.28	**	-	-
PLACE50000022	44.41	69.47	62.06	68.78	41.05	40.65	48.9	41.84	44.88	**	-	-
PLACE50000024	5.49	3.54	5.82	3.22	3.37	3.62	2.52	4.61	6.39	*	-	-
PLACE50000036	104.88	76.2	100.83	90.5	90.12	87.43	80.48	55.18	59.68	*	-	-
PLACE50000059	13.88	12.58	10.61	5.28	9.28	8.17	6.41	11.67	8.16	*	-	-
PLACE50000076	102.38	115.08	148.87	71.48	155.48	121.21	73.46	149.18	129.94	*	-	-
PLACE5000117	10.83	17.74	18.15	14.79	20.06	15.63	5.73	9.53	5.57	*	-	-
PLACE5000143	12.99	9.49	15.16	8.13	9.83	8.05	5.81	8.88	8.8	*	-	-
PLACE5000152	183.88	202.8	98.28	147.85	104.4	116.95	74.92	70.73	75.92	*	-	-
PLACE5000154	90.81	108.58	44.13	75.4	77.68	45.59	40.84	23.33	50.85	*	-	-
PLACE5000155	33.2	24.58	28.07	18.94	31.52	23.72	16.87	17.93	27.02	*	-	-
PLACE5000185	113.75	76.52	82.55	90.32	103.03	88.35	32.72	55.94	39.38	*	-	-
SKNMC1000004	1323.5	940.91	1083.3	685.71	1250.5	673.39	1094.0	1184.2	1080.5	*	-	-
SKNMC1000011	8.28	9.08	11.81	4.48	12.47	5.45	4.79	5.44	8.03	*	-	-
SKNMC1000013	14.72	15.01	20.62	10.84	11.72	12.41	4.74	6.54	6.23	**	-	-
SKNMC1000014	20.79	23.82	26.42	20.23	26.11	18.1	14.78	15.14	11.78	**	-	-
SKNMC1000018	9.56	8.87	12.44	5.32	8.73	4.54	3.81	4.41	5.25	**	-	-
SKNMC1000020	25.49	27.02	25.39	13.83	25.33	12.68	18.7	20.72	21.85	**	-	-
SKNMC1000045	24.5	15.38	17.74	13.48	22.27	16.07	12.53	14.97	15.85	*	-	-
SKNMC1000050	8.51	7.09	7.49	5.47	4.73	4.3	4.47	21.02	3.74	**	-	-
SKNMC1000052	28.94	17.98	24.81	14.13	15.7	9.74	23.81	36.47	28.61	*	-	-
SKNMC1000070	11.6	7.44	13.24	7.34	10.78	7.51	4.21	5.3	7.09	*	-	-
SKNMC1000082	22.01	18.02	26.11	8.38	13.04	8.62	12.8	21.17	18.8	*	-	-
SKNMC1000091	38.26	39.39	43.7	12.98	23.22	18.68	24.3	27.28	22.42	**	-	-
SKNMC1000099	11.34	8.86	15.64	6.38	7.5	4.4	5.83	7.16	6.38	*	-	-

145 / 157

2729

2730

SKINC1000104	16.76	17.55	17.05	8.62	14.24	10.52	9.72	10.59	13.24	*	00	-	-
SKINC1000113	45.53	34.91	35.51	20.7	25.14	23.85	18.88	24.14	23.6	*	0	-	-
SKINC1000119	17.56	14.67	21.54	10.33	12.31	8.53	8.83	13.02	8.17	*	0	-	-
SKINC1000142	18.52	20.57	23.49	7.26	16.72	11.75	8.01	11.68	14.09	*	0	-	-
SKINC1000170	242.49	183.41	211.38	201.28	228.18	182.82	247.45	216.7	250.73	*	00	-	-
SKINC1000178	15.01	12.68	14.32	6.9	9.12	6.49	4.97	7.62	9.01	**	00	-	-
SKINC1000184	7.33	8.61	11.61	2.23	3.73	3.19	2.64	4.74	3.45	*	0	-	-
SKINC1000190	18.76	29.87	24.52	7.87	13.1	9.08	5.41	5.23	5.02	*	00	-	-
SKINC1000225	100.44	119.68	87.32	52.94	84.28	50.45	35.91	33.78	44.06	**	00	-	-
SKINC1000249	23.35	22.68	30.9	13.07	19.78	17.10	12.83	14.11	14.19	*	0	-	-
SPLER1000007	10.7	10.7	16.74	6.92	9.27	7.23	6.86	7.37	5.89	*	0	-	-
SPLER1000012	15.31	13.75	18.53	8.29	13.85	10.56	13.22	11.71	11.02	*	00	-	-
SPLER1000014	74.22	59.99	83.53	32.33	49.13	32.58	29.87	29.67	37.64	*	00	-	-
SPLER1000036	14.69	11.54	10.05	7.95	8.19	8.93	3.18	3.78	2.42	*	00	-	-
SPLER1000059	39.71	33.5	32.87	12.1	19.81	12.24	19.97	18.35	17.18	**	00	-	-
SPLER1000088	20.63	22.57	21.07	11.0	16.29	10.46	9.19	11.89	8.79	*	00	-	-
SPLER1000072	19.68	16.93	14.96	6.08	8.86	7.47	5.18	4.49	6.78	**	00	-	-
SPLER1000101	71.97	43.16	57.48	18.69	27.56	26.6	32.73	48.27	19.97	*	00	-	-
SPLER1000108	7.69	6.28	7.07	3.65	4.36	2.87	3.12	2.84	2.64	**	00	-	-
SPLER1000113	11.47	6.64	12.13	7.54	8.23	7.26	6.6	8.27	7.69	*	00	-	-
SPLER1000114	18.01	14.76	14.38	7.29	9.65	6.3	8.14	8.22	6.73	**	00	-	-
SPLER1000132	5.98	4.51	7.18	2.74	4.42	2.94	1.47	2.25	2.21	*	0	-	-
SPLER1000155	10.67	8.18	9.19	3.88	4.71	4.73	4.9	2.38	2.97	**	00	-	-
SPLER1000136	23.91	23.93	20.23	10.77	11.86	10.15	18.69	14.04	13.77	**	00	-	-
SPLER1000141	42.68	39.73	42.83	27.16	32.91	23.4	23.31	18.41	20.57	**	00	-	-
SPLER1000184	15.72	13.33	13.69	7.78	12.46	6.69	9.59	8.11	9.12	**	00	-	-
SPLER1000188	12.98	9.08	11.45	4.56	6.01	5.04	6.49	5.17	6.32	**	0	-	-
SPLER1000175	19.28	16.94	15.6	7.48	9.23	6.92	11.34	13.49	13.41	**	0	-	-
SPLER1000182	67.34	58.35	68.22	30.69	30.43	27.88	28.16	29.1	23.7	**	00	-	-
SPLER1000185	6.38	8.35	6.58	4.69	4.58	3.06	2.88	2.31	1.51	**	00	-	-
THYRU1000004	48.78	34.98	41.97	30.97	32.65	27.64	26.98	17.5	13.12	*	0	-	-
THYRU1000009	14.59	13.55	14.68	8.47	10.1	5	7.34	4.91	6.62	*	00	-	-
THYRU1000015	18.34	18.66	12.68	8.34	11.37	5.93	10.39	10.96	13.03	*	0	-	-
THYRU1000016	6.78	4.3	5.54	2.79	2.61	1.81	5.14	3.5	4.24	*	0	-	-
THYRU1000023	6.35	6.23	6.53	4.39	3.85	4.2	3.1	5.69	3.88	**	00	-	-
THYRU1000034	192.65	192.5	175.97	306.71	379.09	270.69	454.49	361.08	389.66	*	00	-	-
THYRU1000035	8.93	7.8	9.14	5.68	4.86	4.81	6.05	3.93	4.38	**	00	-	-
THYRU1000037	9.69	6.41	7.38	4.28	3.96	3.33	4.24	3.59	3.83	*	0	-	-
THYRU1000042	6.85	3.93	5.82	2.75	3.24	2.02	2.08	1.63	1.09	*	0	-	-
THYRU1000047	25.64	26.03	24.74	13.43	16.94	11.18	26.77	21.68	23.81	**	00	-	-
THYRU1000080	20.17	51.54	27.22	17.42	19.98	26.99	13.08	15.8	14.07	*	0	-	-
THYRU1000094	11.12	6.99	9.51	6.45	6.49	4.35	11.14	8.41	9.31	*	0	-	-
THYRU1000109	228.62	201.78	195.22	185.65	173.29	154.39	235.81	205.84	221.79	*	00	-	-
THYRU1000127	44.2	30.97	37.5	12.05	18.11	17.75	24.49	15.04	18.2	**	0	-	-
THYRU1000130	21.64	17.3	15.84	7.65	12.45	8	10.24	3.41	8.22	*	0	-	-
THYRU1000137	6.81	10.03	8.23	3.35	5.2	3.34	4.41	2.7	3.47	*	0	-	-
THYRU1000146	11.97	7.39	7.48	8.11	7.24	8.17	5.13	3.03	6.19	*	0	-	-
THYRU1000159	10.45	10.21	8.17	8.28	8.42	3.77	4.54	3.62	4.78	*	00	-	-
THYRU1000163	8.08	8.93	9.7	5.33	5.95	3.66	7.57	4.17	4.21	**	00	-	-
THYRU1000167	67.79	61.53	64.02	27.07	31.08	17.27	75.27	48.55	70.31	**	00	-	-
THYRU1000186	8.38	8.15	3.89	3.94	4.49	2.65	3.38	2.91	3.02	*	0	-	-
THYRU1000197	24	17.58	11.71	8.21	11.63	5.25	6.04	5.8	4.42	*	0	-	-
THYRU1000205	18.24	8.69	6.72	5.28	7.77	4.55	4.15	3.45	6.46	*	0	-	-
THYRU1000234	39.83	32.92	35.9	29.82	39.02	28.83	26.88	25.91	22.08	*	0	-	-
THYRU1000235	34.67	27.15	24.29	22.39	36.44	22.35	22.09	18.28	23.65	*	0	-	-
THYRU1000236	16.2	14.28	11.79	9.92	10.8	4.38	7.94	4.75	8.23	*	0	-	-
THYRU1000240	9.47	7.71	9.48	4.32	5.83	3	5.07	3.54	5.01	*	00	-	-
THYRU1000281	18.28	19.76	15.83	8.28	8.85	4.81	6.42	5.28	8.19	**	00	-	-
THYRU1000287	16.85	13.02	12.08	8.02	8.96	4.34	7.22	6.1	5.44	*	00	-	-
THYRU1000270	9.06	13.03	10.58	5.53	9.48	4.69	20.02	18.33	15.63	*	0	-	-
THYRU1000272	21.67	19.82	18.73	19.89	17.94	12.39	9.77	8.74	9.5	**	00	-	-
THYRU1000284	9.01	5.58	5.31	3.9	6.43	2.97	10.41	8.45	10.27	*	0	-	-
THYRU1000285	28.04	38.58	22.78	13.84	17.05	13.37	10.59	14.52	10.12	*	0	-	-
THYRU1000286	13.56	10.42	8.08	7.5	9.36	4.87	8.84	6.5	4.47	*	0	-	-

146 / 157

2731

2732

THYR01000037	51.14	69.3	41.47	28.99	58.4	26.89	61.05	53.08	61.77				
THYR01000038	26.07	28.19	30.84	13.75	13.51	9.94	17.56	21.51	22.34	**	*	-	-
THYR01000039	4.26	2.01	4.16	1.77	3.46	3.25	2.69	3.3	2.41				
THYR01000039	9.62	5.87	9.26	3.68	8.63	4.69	5.9	7.16	4.32				
THYR01000107	70.48	59.74	59.68	39.34	64.57	40.78	35.76	61.82	50.34				
THYR01000111	13.98	11.92	12.38	7.04	9.78	5.89	8.65	11.53	9.68	*	*	-	-
THYR01000121	31.45	32.43	38.13	15.08	16.17	17.71	26.05	28.32	34.79	**		-	-
THYR01000124	54.63	35.35	48.22	37.6	35.37	31.52	34.87	27.99	31.75				
THYR01000129	8.69	9.01	8.49	5.14	9.82	6.96	8.93	8.73	10.55	*		-	-
THYR01000130	12.23	8.25	10.08	6.62	6.95	6.32	6.99	9.76	9.68	*		-	-
THYR01000132	7.93	8.79	12	5.35	7.15	7.04	4.12	4.66	3.86	*	*	-	-
THYR01000134	17.2	8.08	9.3	6.24	6.27	12.5	5.33	7.31	5.41				
THYR01000144	23.68	30.68	32.27	27.55	32.97	28.83	18.04	19.4	19.02	*		-	-
THYR01000155	35.26	37.35	35.3	18.89	24.31	20.42	31.31	40.6	29.66	**		-	-
THYR01000156	37.13	43.79	38.99	24.77	34.58	29.68	18.78	38.39	21.81	*		-	-
THYR01000183	58.18	53.4	58.15	23.96	28.08	23.48	11.7	14.01	16.02	**	**	-	-
THYR01000173	8.52	8.35	6.68	5.25	10.18	4.15	3.25	4.22	4.73	**		-	-
THYR01000186	35.39	31.09	38.48	14.41	13.8	14.41	7.62	12.05	15.91	**	**	-	-
THYR01000187	17.39	15.95	24.62	6.63	8.73	8.87	5.49	7.02	6.9	*	**	-	-
THYR01000190	7.06	7.97	10.31	6.43	6.1	6.9	4.78	5.24	5.17	*		-	-
THYR01000190	8.72	14.15	13.33	14.25	10.93	7.74	3.75	8.65	8.84				
THYR01000197	13.75	17.13	15.02	14.68	17.34	18.5	8.34	11.47	6.38	*		-	-
THYR01000199	6.8	9.71	11.38	6.6	11.32	10.32	5.5	8.51	7.17				
THYR01000206	20.09	21.74	25.04	13.14	17.5	12.03	7.45	8.8	7.63	*	**	-	-
THYR01000221	6.98	7.15	5.44	4.48	8.31	3.84	3.81	4.45	3.68	*		-	-
THYR01000222	58.33	56.17	63.27	37.58	40.7	31.29	35.13	33.5	43.65	**	**	-	-
THYR01000226	148.64	127.09	184.25	73.36	64.55	57.59	55.08	55.21	55.09	**	**	-	-
THYR01000241	7.79	5.72	9.15	4.42	5	4.49	2.68	4.65	4.26	*	*	-	-
THYR01000242	18.18	23.99	29.76	25.09	51.85	27.45	12.5	20.02	16.81				
THYR01000246	21.67	18.12	24.56	19.46	20.13	15.33	9.29	10.17	9.72	**		-	-
THYR01000253	10.77	13.3	16.4	9.6	16.01	8.45	8.71	8.04	5.73	*		-	-
THYR01000270	12.58	12.99	17.24	8.11	13.02	8.87	9.08	8.16	7.99	*		-	-
THYR01000279	15.04	13.63	18	8.28	14.2	7.73	7.71	7.11	4.92	**		-	-
THYR01000285	1215.2	1221.3	1148.5	961.84	1121.8	818.57	1261.8	1125.7	1518.7				
THYR01000288	24.11	22.65	22.01	13.89	25.53	14.41	22.83	21.93	20.34				
THYR01000296	40.86	36.2	38.51	33.51	37.84	29.24	32.78	49.02	40.29				
THYR01000320	12.29	9.4	14.15	6.59	7.17	6.08	5.28	11.64	9.29	*		-	-
THYR01000322	37.49	34.76	35.09	17.69	26.05	19.02	19.43	23.34	20.74	**	**	-	-
THYR01000327	29.25	26.35	18.48	16.99	27.29	15.75	10.07	12.24	13.62	*		-	-
THYR01000343	16.65	12.58	17.82	8.18	11.42	7.21	10.1	8.05	8.44	*	*	-	-
THYR01000345	6.25	7.77	6.06	4.74	8.59	3.02	4.68	4.72	4.12	*	*	-	-
THYR01000358	13.06	10.33	11.35	6.24	8.78	7.32	8.15	6.51	9.21	*	*	-	-
THYR01000368	5.38	5.26	6.26	3.15	3.94	2.68	6.19	3.05	2.43	**		-	-
THYR01000375	14.1	11.6	18.64	8.68	5.99	5.1	3.89	3.2	3.66	*	**	-	-
THYR01000381	10.31	8.3	9.78	4.37	5.3	5.86	4.17	5.35	3.69	**	**	-	-
THYR01000387	11.39	10.61	13.54	4.54	5.97	3.48	10.69	6.84	3.97	**		-	-
THYR01000394	61.78	50.97	22.02	14.87	28.91	8.08	6.98	9.73	11.33	*	*	-	-
THYR01000395	29.06	37.87	27.54	18.13	24.45	14.15	25.79	23.19	25.7	*		-	-
THYR01000400	10.51	10.51	9.69	3.23	12.01	6.83	3.72	3.73	3.08	**		-	-
THYR01000401	10.42	9.53	10.77	8.74	8.08	4.73	3.24	5.18	5.6	**	**	-	-
THYR01000407	111.95	92.99	98.17	83.29	76.05	58.17	51.33	55.33	55.02	**	**	-	-
THYR01000420	15.01	13.25	17.92	11.1	11.82	8.58	12.79	12.37	14.41	*		-	-
THYR01000438	8.38	5.22	8.85	5.12	4.11	4.31	4.22	4.7	3.13	*		-	-
THYR01000452	16.04	12.59	12.87	5.92	10.6	6.13	2.16	3.93	4.57	*	**	-	-
THYR01000455	7.17	8.57	9.52	2.62	3.31	3.44	2.83	2.18	2.11	**	**	-	-
THYR01000471	78.16	61.99	62.01	65.6	62.48	70.88	61.77	51.28	74.08				
THYR01000481	8.9	7.89	8.46	4.91	7.54	4.12	2.78	3.16	5.02	**		-	-
THYR01000484	19.83	12.78	16.87	10.37	11.59	9.38	11.21	9.7	11.81	*		-	-
THYR01000488	28.64	20.75	23.35	20.89	21.13	19.69	15.6	20.62	19.42				
THYR01000501	11.59	12.72	13.28	7.74	9.03	6.79	8.05	7.95	10.26	**	*	-	-
THYR01000502	55.88	62.65	49.31	44.97	46.78	38.57	44.58	40.93	38.11	*	*	-	-
THYR01000505	9.64	7.65	7.27	5.58	5.82	3.88	3.98	5.47	2.02	*	*	-	-
THYR01000535	37.32	41.34	38.41	38.72	38.24	31.02	23.49	20.01	18.32	**		-	-
THYR01000566	102.92	92.57	85.3	76.31	104.96	82.73	51.58	50.77	49.93	**		-	-

147 / 157

2733

2734

THYR01000554	8.64	7.17	8.38	4.16	8.61	9.01	4.28	2.04	3.45	*	-
THYR01000560	8.63	4.82	4.79	4.77	4	3.41	3.58	3.02	3.94	*	-
THYR01000570	9.09	8.96	8.83	5.27	8.38	8.54	6.25	4.89	7.09	*	-
THYR01000572	52.16	31.62	39.78	12.82	33.15	30.73	30.68	23.84	11.23	*	-
THYR01000573	8.86	8.49	8.82	4.33	8.49	8.35	13.45	11.85	8.38	*	-
THYR01000577	9.73	7.83	7.69	3.83	6.26	2.6	2.58	3.82	2.74	*	-
THYR01000580	40.56	32.54	29.85	22.22	19.88	19.81	24.19	17.34	26.62	*	-
THYR01000584	39.86	36.8	31.89	19.25	26.6	16.24	28.15	21.02	19.13	*	-
THYR01000585	8.89	7.37	8.85	4.19	8.67	4.88	5.4	3.58	2.99	*	-
THYR01000588	43.85	33.14	32.61	22.84	31.85	11.59	15.52	10.88	14.47	*	-
THYR01000802	18.04	15.25	10.78	7.23	11.51	11.78	8.67	7.72	10.81	*	-
THYR01000809	10.08	9.44	5.91	8.29	6.99	4.61	4.9	5.21	5.08	*	-
THYR01000815	134.18	117.87	85.87	88.81	98.31	67.18	77.11	87	84.37	*	-
THYR01000825	33.27	24.89	18.68	12.6	10.75	20.13	8.18	8.88	11.76	*	-
THYR01000838	17.91	18.49	14.48	12.48	11.49	8.7	8.72	6.17	8.59	*	-
THYR01000837	18.22	22.2	13.43	10.39	14.05	9.78	9.07	5.2	7.28	*	-
THYR01000841	7.23	5.87	6.32	4.82	10.06	3.87	8.42	4.55	2.07	*	-
THYR01000857	10.91	9.88	8.65	8.93	7.74	7.68	9.53	8.35	10.94	*	-
THYR01000858	14.37	9.48	5.1	6.28	5.89	5.43	4.59	4.82	4.77	*	-
THYR01000862	10.63	9.43	8.48	6.7	5.85	6.96	7.92	7.09	10.89	*	-
THYR01000866	30.51	33.5	45.43	12.9	20.38	13.81	14.6	13.53	16.48	*	-
THYR01000876	11.56	11.35	8.48	6.32	9.3	8.87	4.16	5.02	6.44	*	-
THYR01000878	23.44	18.73	20.73	10.13	15.99	8.49	9	7.18	9.87	*	-
THYR01000884	27.78	27.85	20.27	12.26	18.45	11.08	26.97	17.82	24.98	*	-
THYR01000894	16.87	11.78	10.72	5.47	10.25	5.89	12.29	9.36	8.6	*	-
THYR01000899	8.11	6.38	4.55	5.62	6.31	3.48	7.37	3.08	5.08	*	-
THYR01000712	17.56	14.87	12.31	9.08	11.99	7.67	12.79	9.26	13	*	-
THYR01000715	29.82	23.26	21.02	20.98	27.37	20.44	15.76	18.98	18.68	*	-
THYR01000718	13.12	15.14	11.94	8.87	12	7.65	7.48	6.69	8.98	*	-
THYR01000717	11.89	15.21	12.45	6.9	9.22	5.08	30.93	6.96	7.18	*	-
THYR01000723	30.36	26.4	21.78	10.82	20.63	14.48	30.94	22.3	30.26	*	-
THYR01000734	18.61	19.81	18.12	11.94	17.61	9.39	9.11	8.61	13.1	*	-
THYR01000748	17.48	14.08	15.67	5	7.52	5.42	7.82	5.84	7.09	*	-
THYR01000756	20.17	21.24	19.68	13.28	17.61	14.04	14.7	23.04	21.01	*	-
THYR01000759	389.81	334.57	369.89	288.51	345.41	273.28	289.37	368.07	273.4	*	-
THYR01000776	4.97	4.89	7.55	4.25	6.16	4.82	2.58	5.02	3.55	*	-
THYR01000777	5.42	5.83	6.34	5.34	6.01	4.03	3.82	3.75	3.21	*	-
THYR01000778	483.37	458.9	483.51	453.87	549.42	449.1	368.6	510.93	458.47	*	-
THYR01000782	17.27	20.25	23.35	14.58	17.12	13.37	15.81	16.4	16.35	*	-
THYR01000783	5.74	5.45	6.68	5.43	9.62	4.88	3.62	5.19	3.8	*	-
THYR01000785	26.82	30.48	26.26	16.78	13.18	14.48	32.89	30.78	26.94	*	-
THYR01000787	50.52	37.21	54.05	28.71	18.06	26.25	20.11	24.5	22.68	*	-
THYR01000792	14.11	12.1	15.65	4.55	8.72	7.1	8.13	7.14	8.33	*	-
THYR01000793	36.92	35.83	42.61	19.08	27.18	19.55	34.91	38.74	32.64	*	-
THYR01000795	37.33	45.29	88.22	42.19	67.38	40.15	12.57	21.17	13.38	*	-
THYR01000796	10.25	11.83	14.01	6.19	10.21	5.98	7	11.88	11.1	*	-
THYR01000798	9.87	14.22	12.83	9.08	8.1	7.49	5.04	7.33	5.2	*	-
THYR01000800	37.89	64.75	19.73	33.2	22.38	23.21	13.79	16.89	13.75	*	-
THYR01000805	8.27	6.21	8.02	5.33	8.39	4.44	3.81	4.14	3.82	*	-
THYR01000815	109.41	82.13	80.82	43.61	38.14	26.61	31.78	21.16	31.98	*	-
THYR01000820	33.88	25.13	34.71	19.68	21.88	18.45	16.87	14.34	12.65	*	-
THYR01000835	9.51	9.82	10.3	6.71	6.28	4.65	5.41	5.24	6.07	*	-
THYR01000843	27.06	18.5	24.95	16.28	19.34	15.73	5.22	7.81	8.06	*	-
THYR01000846	7.88	5.91	13.48	4.53	6.28	7.74	3.89	7.02	5.28	*	-
THYR01000852	24.68	25.37	23.86	12.87	20.2	12.5	9.33	18.39	11.62	*	-
THYR01000855	7.12	7.47	10.43	2.52	5.79	2.98	4.3	5.1	5.2	*	-
THYR01000855	9.21	5.67	8.84	3.4	8.58	3.18	5.99	8.76	7.67	*	-
THYR01000886	57.27	48.1	51.89	32.85	43.74	27.87	46.24	38.05	41.55	*	-
THYR01000881	13.67	12.21	13.18	9.6	10.62	12.24	10.21	18.82	10.81	*	-
THYR01000894	9.33	7.11	11.4	5.68	6.12	4.31	4.14	4.68	4.75	*	-
THYR01000895	8.9	5.89	10.11	3.35	4.07	4.72	4.32	4.09	3.45	*	-
THYR01000916	9.19	7.72	18.87	5.02	8.87	5.27	4.16	8.12	3.42	*	-
THYR01000917	6.89	10.63	12.32	4.97	9.61	5.39	5.77	7.94	8.2	*	-
THYR01000926	14.99	10.81	12.37	6.78	9.85	8.06	7.42	7.03	11.27	*	-

148 / 157

2735

2736

THYRO1000934	9.21	6.94	9.67	4.17	7.52	4.26	6.41	5.73	4.96	*	*	-	-
THYRO1000951	21.19	13.82	17.24	10.15	10.71	6.54	7.11	11.34	11.56	*	*	-	-
THYRO1000952	38.82	45.17	51.97	24.89	24.37	18.77	17.55	26.97	13.34	**	**	-	-
THYRO1000956	11.42	8.41	13.85	5.27	6.34	6.15	5.07	6.61	6.85	*	*	-	-
THYRO1000960	9.97	3.83	6.63	2.78	4.13	2.34	2.46	2.03	6.43	*	*	-	-
THYRO1000961	33.04	35.26	36.48	15.01	15.48	16.21	21.6	24.41	21.14	**	**	-	-
THYRO1000964	14.43	12.46	17.93	3.92	6.7	5.82	5.43	7.53	4.68	**	**	-	-
THYRO1000971	15.18	18.5	18.14	8.34	13.54	9	9.38	6.59	10.62	*	**	-	-
THYRO1000974	7.51	5.29	9.43	2.24	6.94	1.8	3.55	5.59	3.45	*	*	-	-
THYRO1000975	7.71	4.75	6.93	3.38	5.71	4.79	3.77	4.87	3	*	*	-	-
THYRO1000983	8.4	4.14	7.44	3.88	5.03	3.36	3.1	3.21	4.03	*	*	-	-
THYRO1000984	9.36	6.4	11.52	6.49	7.32	5.22	5.08	2.55	2.85	*	**	-	-
THYRO1000988	10.2	6.67	7.82	4.27	4.23	4.8	2.94	2.66	2.92	*	**	-	-
THYRO1000991	12.33	10.95	12.47	5.27	7.49	7.34	5.57	6.15	6.01	**	**	-	-
THYRO1000999	9.77	12.84	10.64	5.58	8.33	5.82	5.68	3.11	2.9	*	**	-	-
THYRO1001003	324.55	449.5	824.63	353.93	491.59	377.12	509.02	394.65	380.17	*	*	-	-
THYRO1001015	16.58	13.37	11.66	7.38	13.81	8.4	12.04	8.04	10.47	*	*	-	-
THYRO1001016	34.7	29.76	32.31	14.48	17.7	10.51	27.97	31.88	30.58	**	*	-	-
THYRO1001022	27.6	17.2	20.61	6.98	9.39	10.85	5.35	8.58	6.8	*	**	-	-
THYRO1001031	16.75	9.57	11.29	5.12	7.05	5.89	8.13	8.39	9.34	*	*	-	-
THYRO1001033	11.34	7.67	8.68	7.55	6.62	6.01	4.3	4.01	4.9	*	*	-	-
THYRO1001062	12.88	12.45	12.49	4.68	5.59	4.88	3.54	3.68	2.79	**	**	-	-
THYRO1001063	10.18	8.38	10.08	5.14	7.17	4.58	3.55	4.32	2.35	*	**	-	-
THYRO1001071	18.49	15.5	18.23	9.12	10.91	7.86	5.73	5.41	5.84	**	**	-	-
THYRO1001080	11.77	12.06	11.47	3.85	12.27	3.17	3.08	4.59	4.41	**	*	-	-
THYRO1001093	12.17	9.63	9.75	5.58	4.91	5	8.58	8.31	7.24	**	*	-	-
THYRO1001100	33.06	27.59	28.02	30.29	27.98	30.28	66.02	40.22	70.38	*	*	-	-
THYRO1001102	18.18	8.7	5.95	5.04	6.9	4.52	4.36	4.71	4.45	*	*	-	-
THYRO1001104	5.1	2.75	2.78	0.69	0.92	1.62	1.32	0.94	0.7	*	*	-	-
THYRO1001109	19.62	18.07	14.28	10.98	10.98	9.78	8.55	11.06	8.68	*	*	-	-
THYRO1001112	18.89	6.97	7.59	2.84	3.69	2.77	2.56	1.4	2.3	*	**	-	-
THYRO1001120	9.27	9	8.82	3.48	4.66	2.88	4.02	3.07	2	**	*	-	-
THYRO1001121	20.41	16.34	16.91	9.1	13.05	6.93	13.22	15.3	13.97	*	*	-	-
THYRO1001128	25.09	18.24	17.74	8.53	9.61	6.74	3.45	4.7	7.82	**	**	-	-
THYRO1001133	7.42	8.4	5.88	4	5.05	4.23	3.88	4.14	3.81	*	**	-	-
THYRO1001134	29.23	23.86	23.28	16.51	20.53	18.49	22.14	16.92	20.47	*	*	-	-
THYRO1001142	5.04	4.99	5.1	2.78	3.05	3.47	2.44	2.28	1.6	**	**	-	-
THYRO1001172	31.85	26.04	28.94	20.38	18.72	13.59	18.5	15.33	18.17	*	**	-	-
THYRO1001175	95.55	98.04	90.56	104.29	114.18	77.1	51.33	48.58	49.17	**	*	-	-
THYRO1001177	10.9	11.18	11.71	5.13	7.17	5.9	4.63	5.82	5.02	**	**	-	-
THYRO1001189	16.52	14.55	11.53	5.76	11.24	3.42	11.07	10.9	8.88	*	*	-	-
THYRO1001194	11.83	6.2	11.19	5.29	7.22	4.28	5.84	7.01	6.86	*	*	-	-
THYRO1001204	62.04	33.32	32.8	18.53	22.19	18.27	39.75	37.94	48.59	*	*	-	-
THYRO1001205	8.98	12.23	9.39	5.39	8.08	5.04	6.55	6.04	4.85	*	*	-	-
THYRO1001213	41.54	37.73	35.88	23.38	25.7	20.79	37.87	32.41	38.52	**	*	-	-
THYRO1001224	10.88	11.43	10.75	6.38	8.69	6.7	7.21	4.53	6.38	*	**	-	-
THYRO1001237	18.76	20.25	15.97	16.59	13.69	5.38	9.78	5.89	6.27	**	*	-	-
THYRO1001242	13.1	19.03	16.08	12.27	8.61	6.73	9.21	5.95	9.02	*	*	-	-
THYRO1001250	27.32	21.68	20.65	8.74	18.76	12.77	10.51	8.88	13.04	*	**	-	-
THYRO1001252	8.57	4.74	4.68	3.78	5.55	4.02	4.79	3.93	5.05	*	*	-	-
THYRO1001256	25.46	17.38	17.48	9.98	11.28	8.76	13.97	15.24	19.55	*	*	-	-
THYRO1001271	27.85	25.6	17.14	13.77	18.43	9.74	22.54	18.92	23.05	*	*	-	-
THYRO1001287	126.74	172.83	126.71	56.78	189.06	71.99	103.16	93.81	120.13	*	*	-	-
THYRO1001290	11.3	8.82	9.68	4.97	7.48	6.55	3.85	5.84	8.41	*	*	-	-
THYRO1001291	13.9	16.44	13.22	8.85	8.33	3.9	6.83	3.84	6.66	*	**	-	-
THYRO1001297	13.17	12.27	11.41	6.73	6.89	4.23	7.87	5.3	6.04	**	**	-	-
THYRO1001302	28.98	24.47	17.01	9.99	19.11	8.25	41.38	28.42	38.98	*	*	-	-
THYRO1001313	3.4	2.28	2.66	1.62	1.97	1.71	3.09	2.85	2.3	*	*	-	-
THYRO1001320	4.85	2.3	4.45	3.81	2.95	2.21	2.87	2.22	1.94	*	*	-	-
THYRO1001321	84.54	108.43	89.51	45.59	48.9	38.67	60.51	58.13	42.33	**	*	-	-
THYRO1001322	15.07	13.59	11.8	5.6	6.9	4.45	5.91	6.73	6.71	**	**	-	-
THYRO1001327	81.14	82.79	47.58	27.21	38.08	29.25	40.32	44.88	30.36	**	*	-	-
THYRO1001336	21.04	18.95	13.64	17.33	15.44	11.88	17.21	14.24	12.35	*	*	-	-
THYRO1001347	42.47	35.49	29.58	20.44	37.43	12.7	20.28	16.34	15.93	**	*	-	-

149 / 157

2737

2738

THYR01001350	6.88	7.77	8.62	4.41	8.39	9.78	5.37	5.5	4.22	0	-
THYR01001351	35.77	35.26	35.97	12.88	20.63	14.65	24.19	31.52	25.12	**	-
THYR01001355	2.39	1.31	4.42	2.3	2.69	1.43	1.82	2.18	1.5	0	-
THYR01001374	10.11	11.14	11.66	5.19	7.75	3.64	2.84	5.3	4.44	*	-
THYR01001401	11.45	8.59	8.77	5.02	5.3	5.96	5.5	8.66	6.31	*	-
THYR01001403	3.12	3.82	7.88	1.61	2.43	2.61	3.73	1.98	3.81	*	-
THYR01001405	84.82	60.76	92.35	68.75	78.6	50.19	53.53	29.8	47.84	*	-
THYR01001406	13.59	7.57	12.02	4.78	8.87	1.76	11.89	12.59	9.54	*	-
THYR01001411	4.72	4.15	5.65	3.88	2.4	3.38	2.62	2.5	3.6	*	-
THYR01001420	9.34	4.82	8.71	4.7	4.71	5.55	3.24	4.39	8.81	*	-
THYR01001426	9.28	4.52	12.27	5.65	6.22	4.37	4.27	8.59	6.13	*	-
THYR01001430	10.74	13.91	18.18	13.99	18.28	13.49	6.82	9.48	10.77	*	-
THYR01001434	7.68	8.88	12.81	8.64	10.68	8.13	5.38	8.91	2.99	*	-
THYR01001456	26.48	36.99	36.98	17.02	23.99	18.02	30.71	34.81	38.85	*	-
THYR01001457	14.33	10.91	13.28	5.72	4.64	3.33	4.05	3.22	4.83	**	-
THYR01001458	18.32	13.83	12.78	10.99	10.13	5.72	4.4	7.29	6.87	*	-
THYR01001459	5.78	4.92	7.15	2.81	3.48	4.24	3.11	4.43	5.61	*	-
THYR01001471	13.48	10.81	13.3	5.13	9.82	6.19	6.36	8.85	9.45	*	-
THYR01001478	8.04	3.79	8.81	2.94	2.62	2.21	4.12	4.22	2.59	*	-
THYR01001480	18.07	18.86	14.48	8.78	13.68	13.34	3.54	8.81	18.35	*	-
THYR01001481	18.23	13.97	17.19	13.3	16.2	13.73	4.94	8.01	3.76	**	-
THYR01001487	39.34	43.94	45.87	41.48	42.62	34.09	36.78	38.23	41.44	*	-
THYR01001496	6.21	3.33	12.71	2.18	5.51	4.67	3.43	3.46	4.85	*	-
THYR01001498	13.62	15.45	16.81	11.32	29.15	8.37	4.81	8.08	4.31	**	-
THYR01001510	9.97	6.87	19.31	8.41	6.21	3.55	4.99	6.45	19.57	*	-
THYR01001512	11.51	8.21	10.34	5.87	6.36	6.84	4.43	18.67	5.22	*	-
THYR01001519	12.05	18.47	10.9	4.73	6.89	4.14	4.42	4.28	6.04	*	-
THYR01001522	18.38	8.84	12.83	8.15	7.71	8	5.13	5.33	7.66	*	-
THYR01001523	15.14	10.51	16.01	7.01	9.2	6.38	3.77	3.19	4.5	*	-
THYR01001526	5.87	5.48	12.84	2.85	6.19	3.09	4.51	3.63	4.01	*	-
THYR01001529	78.62	99.68	81.07	73.62	81.82	35.68	29.07	23.9	34.89	*	-
THYR01001534	18.48	7.92	11.72	8.09	7.42	11.24	8.53	6.35	3.17	*	-
THYR01001537	5.9	3.05	5.93	1.4	2.75	2.75	3.28	2.87	4.81	*	-
THYR01001541	27.84	38.72	38.42	23.51	30.22	18.99	22.01	26.98	14.85	*	-
THYR01001545	12.87	9.78	11.61	3.68	6.9	5.41	4.16	22.14	4.44	**	-
THYR01001559	143.87	217.09	127.43	126.92	211.99	217.97	165.83	143.91	101.7	**	-
THYR01001563	29.68	25.68	28.83	14.46	17.77	19.17	10.82	18.32	17.35	**	-
THYR01001570	5.85	5.24	10.72	3.21	3.79	2.31	2.47	4.03	3.05	*	-
THYR01001573	19.58	14.52	16.67	7.88	10.88	10.26	4.93	9.17	5.78	*	-
THYR01001584	11.71	9.79	13.05	7.76	12.43	7.3	8.19	4.92	4.88	*	-
THYR01001593	119.63	87.93	122.57	37.27	54.3	33.58	100.89	84.41	118.61	**	-
THYR01001595	9.11	9.13	10.8	7.84	8.88	8.77	4.62	12.97	7.82	*	-
THYR01001596	8.25	3.54	8.97	3.03	4.5	2.76	3.54	3.67	4.02	*	-
THYR01001602	7.1	6.21	7.8	6.53	4.98	5.02	4.21	4.12	2.97	**	-
THYR01001605	28.22	39.54	25.87	13.48	23.89	22.57	11.83	10.02	13.09	*	-
THYR01001608	19.26	17.94	17.77	8.3	22	8.49	8.07	8.26	11.18	**	-
THYR01001617	7.38	9.07	12.28	4.4	5.61	3.63	3.83	3.37	4.51	*	-
THYR01001634	9.8	9.15	14.08	5.85	9.76	4.62	5.62	3.69	4.66	*	-
THYR01001637	6.56	2.42	6.39	2.18	3.8	2.31	2.41	3.7	6.48	*	-
THYR01001641	12.87	8.05	12.44	6.03	4.87	6.49	3.32	2.86	2.92	*	-
THYR01001656	28.03	12.66	30.89	7	10.07	5.22	6.32	7.53	5.64	*	-
THYR01001658	7.58	6.85	10.14	4.85	4.77	3.94	3.38	2.8	2.5	*	-
THYR01001661	150.1	98.3	106.51	135.25	186.2	125.8	74	80.32	53.76	*	-
THYR01001671	10.97	8.43	12.21	5	5.04	3.97	5.12	5.12	3.84	**	-
THYR01001672	12.9	11.31	11.75	5.85	6.91	5.08	3.89	2.74	2.72	**	-
THYR01001673	31.04	28.57	28.79	16.85	18.7	19.9	22.8	23.43	23.51	**	-
THYR01001677	8.63	8.88	9.87	3.55	4.9	4.2	6.4	8.55	8.85	*	-
THYR01001683	38.34	16.85	25.87	16.11	18.7	19.17	8.36	10.78	9.09	*	-
THYR01001700	12.31	7.03	9.15	3.61	4.23	3.04	2.79	3.83	4.8	*	-
THYR01001702	23.7	21.62	17.4	10.57	6.8	11.44	6.01	4.83	2.94	*	-
THYR01001703	7.22	4.23	6.14	2.63	3.23	1.81	2.37	2.42	2.04	*	-
THYR01001708	8.39	5.62	8.09	3.72	3.11	3.37	3.24	2.65	1.28	*	-
THYR01001721	11.69	10.42	16.73	6.09	4.96	4.48	3.92	2.33	3.87	*	-
THYR01001725	147.69	144.4	144.74	111.99	128.3	88.63	130.43	108.81	93.76	*	-

150 / 157

2740

THYR61001730	7.74	3.57	7.41	4.48	4.19	3.07	3.84	4.09	4.72		
THYR61001738	1.57	3.03	2.24	0.97	0.17	0.58	0.7	0.99	1.64	*	
THYR61001743	5.12	3.48	3.39	1.87	2.12	2.94	1.98	2.8	2.48		
THYR61001745	30.25	14.84	20.8	6.81	4.85	9.25	12.7	8.09	11.05	*	
THYR61001746	18.78	5.89	7.69	2.83	3.1	3.64	3.64	1.32	2.13	*	*
THYR61001770	28.01	12.02	16.57	10.05	13	10.49	5.37	6.07	5.43	*	*
THYR61001772	29.2	58.08	20.11	28.47	34.34	15.95	9.21	8.24	9.37	*	*
THYR61001778	39.03	35.61	22.42	13.39	32.45	16.04	9.42	8.03	11.2	*	*
THYR61001783	9.35	3.26	5.88	4.02	4.49	3.37	4.45	4.28	5.77	*	*
THYR61001786	66.58	52.34	52.35	31.51	31.58	30.08	74.83	82.87	78.59	**	
THYR61001800	25.21	17.42	21.74	18.92	18.43	13.71	8.41	10.02	11.59	*	**
THYR61001803	12.96	11.11	10.61	7.04	8.3	3.44	5.71	5.48	8.84	*	**
THYR61001809	19.44	7.96	8.8	4.59	3.72	3.71	3.82	1.76	1.34	*	**
THYR61001817	18.78	13.28	15.51	15.93	13.63	6.93	7.89	4.88	8.37	*	*
THYR61001819	14.15	14.31	9.69	5.8	6.03	6.06	6.27	4.29	5.81	*	*
THYR61001828	8.14	8.22	6.85	5.42	7.94	4.29	7.29	4.28	2.47	*	*
THYR61001854	50.53	32.68	37.71	33.17	34.18	22.81	22.18	27.13	21.15	*	*
THYR61001895	9.88	7.83	3.92	4.41	4.63	2.55	3.7	10.54	3.48		
THYR61001907	27.03	13.41	12.49	9.48	10.64	7.02	4.92	4.41	6.47		
TRACH1000096	5.44	7.16	8.08	5.68	4.68	5.63	6.46	4.52	6.78	*	
TRACH1000133	9.02	5.8	8.2	2.58	4.45	3.1	3.07	2.02	4.14	*	*
TRACH1000074	13.69	8.32	8.63	4.56	7.68	4.06	6.37	3.07	3.63	*	*
TRACH1000095	11.28	8.48	10.55	5.84	3.9	3.41	3.79	4.56	3.68	**	*
TRACH1000102	28.88	26.3	20.1	15.81	18.57	15.2	32.52	24.6	27.09	*	*
TRACH1000158	84.66	69.61	40.48	28.57	26.39	20.8	59.3	56.03	66.91	*	*
TRACH1000126	17.36	13.19	13.78	8.97	9.6	6.46	5.91	5.75	8.51	*	**
TRACH1000145	7.25	6.83	5.07	5.29	3.75	3.17	3.49	3.25	4.07	*	*
TRACH1000180	11.76	11.24	7.7	4.95	6.58	4.95	5.32	4.22	5.87	*	*
TRACH1000184	33.51	45.84	50.29	14.82	25.81	22.15	25.29	16.88	23.76	*	*
YESER1000064	11.35	8.9	8.13	6.46	9.69	6.32	5.53	5.44	6.01	*	*
YESER1000087	32.64	30.13	26.81	16.7	21.99	18.54	37.52	29.73	33.83	**	*
YESER1000013	13.16	11.68	12.05	7.19	8.9	4.25	7.18	4.46	8.92	*	*
YESER1000029	23.17	20.53	28.94	10.16	15.29	14.78	9.13	12.72	8.48	*	**
YESER1000059	4.76	2.32	6.57	3.56	3.32	3.96	1.16	3.33	1.98		
YESER1000100	8.85	5.84	8.92	5.95	6.82	5.48	3.89	5.59	3.98	*	
YESER1000107	185.28	125.83	154.47	126.48	135.37	133.6	74.61	79.36	82.66	**	*
YESER1000117	3.74	4.02	10.93	3.48	5.02	4.1	2.18	2.11	2.69		
YESER1000122	5.84	5.31	7.17	3.91	4.47	3.29	2.52	3.75	2.41		
YESER1000137	7.93	10.91	7.7	8.31	5.46	4.02	3.34	3.30	6.77	*	
YESER1000195	12.47	4.76	10.7	4.43	4.22	9.07	3.11	4.32	13.25		
YESER1000215	89.85	69.16	78.83	74.48	93.71	81.06	50.08	36.95	45.24		
YESER1000279	16.42	11.12	19.97	7.88	19.23	5.52	8.64	13.68	21.33		
YESER1000353	40.43	28.46	56.48	41.4	53.72	50.13	17.51	33.82	20.25		
YESER1000388	4.51	6.13	8.09	5.58	8.45	6.21	3.44	6.29	2.57		
YESER1000394	137.33	122.92	135.03	107.32	177.13	115.58	81.85	76.56	86.04	**	*
YESER1000410	7.84	11.73	12.31	7.37	8.65	7.32	5.82	6.16	5.5	*	*
YESER1000411	14.34	16.7	11.87	14.26	14.7	9.53	8.54	4.28	7.07	**	*
YESER1000415	6.65	6.22	8.5	3.18	5.01	4.35	2.06	3.93	7.37	*	
YESER1000440	7.1	4.56	7.43	10.06	4.89	9.04	3.62	6.97	6.76		
YESER1000452	127.08	114.54	102.58	100.33	108.23	122.79	77.87	58.16	82.71	*	
YESER1000539	57.4	47.68	67.32	38.3	45.7	42.57	51.56	51.15	50.18		
YESER1000554	7.29	12.84	14.13	6.95	4.57	6.75	3.69	3.4	3.85	*	
YESER1000557	4.5	6.91	10.86	5.84	5.73	4.35	3.05	2.98	4.78		
YESER1000575	16	18.27	24.28	11.3	16.07	11.77	13.59	13.83	16.27		
YESER1000585	14.4	14.28	13.88	9.87	13.37	8.72	4.91	4.08	5.01	**	*
YESER1000592	28.09	20.04	22.75	8.88	11.63	7	7.53	8.48	8.84	**	*
YESER1000596	8.27	6.98	6.99	4.92	3.93	3.11	3.31	12.18	8.08	*	*
YESER1000688	7.65	11.62	18.09	8.2	9.11	7.91	4.21	8.98	3.61		
YESER1000743	27.12	25.27	28.49	19.46	23.17	18.77	12.01	15.14	12.53	**	*
YESER1000752	23.78	18.91	24.81	11.19	11.63	9.8	14.24	18.48	18.19	*	*
YESER1000781	7.03	8.3	9.9	5.84	5.41	3.65	3.1	8.25	5.47	*	*
YESER2000039	16.58	20.4	16.46	7.12	12.23	7.45	6.23	4.66	8.91	*	*
YESER2000102	15.76	17.97	15.75	7.45	12.62	8.13	11.35	8.16	13.31	*	*
YESER2000164	69.66	54.85	54.07	10.32	20.39	15.6	15.69	21.61	20.97	**	**

2741

2742

YESER2000175	7.45	8.05	7.39	3.43	6.1	3.72	4.54	8.1	4.35	*	-	-
YESER2000180	18.04	20.73	21.8	11.19	14.77	10.58	8.74	12.64	12.61	**	-	-
YESER2000180	11.71	8.48	9.88	8.59	8.24	5.52	6.89	8.28	14.79	*	-	-
YESER2000200	150.21	142.83	144.63	172.84	215.31	128.05	93.38	87.48	97.05	**	-	-
YESER2000204	7.05	7.07	9.65	4.56	4.24	3	2.19	4.63	3.19	*	-	-
YESER2000218	16.66	12.07	17.29	7.31	8.24	8.18	8.32	5	5.15	**	-	-
YESER2000230	12.09	13.48	13.57	4.57	10.45	4.08	3.93	4.19	2.98	*	-	-
YESER2000272	8.95	9.01	11.14	4.32	3.68	3.22	4.38	3.53	4.22	**	-	-
YESER2000299	8.61	7.23	11.81	5.04	5.83	4.25	3.6	5.71	4.76	*	-	-
YESER2000323	17.3	12.74	15.22	15.38	17.47	14.24	5.84	8.46	4.65	**	-	-
YESER2000327	18.5	9.19	14.45	6.81	8.17	7.45	3.2	5.07	5.85	*	-	-
YESER2000328	7.53	7.78	8.38	3.45	4.17	4.12	2.35	2.19	1.81	**	-	-
YESER2000330	25.07	20.15	22.73	13.34	18.45	13.18	15.83	15.15	15	*	-	-
YESER2000336	7.29	9.37	11.82	5.88	8.09	5.78	3.84	3.5	5	*	-	-
YESER2000354	10.99	9.78	10.4	5.27	8.22	4.93	3.58	3.58	2.59	*	-	-
YESER2000378	188.58	110.88	139.59	81.44	99.69	74.23	74.17	76.59	54.75	*	-	-
YESER2000379	32.08	21.47	28.34	16.72	15.12	13	22.77	26.45	21.84	*	-	-
YESER2000397	21.17	10.11	25.4	12.77	12.2	10.8	15.72	16.01	17.99	*	-	-
YESER2000418	5.32	5.21	7.48	3.5	6.28	4.01	2.87	3.84	3.48	*	-	-
YESER2000420	8.47	4.35	7.18	4.04	3.81	2.46	2.31	4.27	1.92	*	-	-
YESER2000430	8.74	5.78	7.33	2.94	5.32	2.11	1.8	2.27	0.71	*	-	-
YESER2000448	14.67	14.58	16.08	4.71	4.34	2.88	3.91	2.29	2.27	**	-	-
YESER2000449	48.1	50.29	39.97	32.76	32.82	21.58	37.62	28.99	23.73	*	-	-
YESER2000458	45.88	32.52	32.47	18.88	21.08	19.47	15.48	22.28	22.31	*	-	-
YESER2000562	6.11	3.13	4.69	5.4	3.95	3.19	3.24	2.45	2.42	*	-	-
YESER2000573	26.67	20.52	18.39	14.29	11.45	8.79	8.94	7.48	12.2	*	-	-
YESER2000604	9.02	7.76	4.59	4.85	4.32	4.42	1.84	4.41	1.34	*	-	-
YESER2000614	9.59	7.78	8.05	4.77	5.05	4.35	3	3.94	2.87	*	-	-
YESER2000630	16.75	10.83	10.74	6.38	9.48	6.28	4.67	4.45	3.5	*	-	-
YESER2000641	15.71	14.29	14.25	7.95	7.6	6.71	8.94	7.41	6.6	**	-	-
YESER2000645	9.28	11.51	10.68	5.27	8.01	4.95	5.42	4.96	6.81	*	-	-
Y79AA1000013	24.68	15.88	15.09	6.08	5.43	5.53	13.59	13.07	18.88	*	-	-
Y79AA1000030	19.52	15.62	14.47	8.07	7.77	10.99	7.86	7.77	10.02	*	-	-
Y79AA1000033	29.91	18.65	28.38	15.37	15.72	11.59	28.15	19.64	28.85	*	-	-
Y79AA1000037	4.94	2.8	4.89	5.26	3.24	0.91	1.44	1.94	2.43	*	-	-
Y79AA1000041	79.88	63.74	64.67	39.77	35.14	29.21	42.77	39.09	45.97	*	-	-
Y79AA1000059	9.8	7.3	8.44	4.37	5.08	2.08	5.32	3.01	5.11	*	-	-
Y79AA1000065	115.14	101.03	108.53	93.49	124.98	111.44	68.74	63.93	85.58	**	-	-
Y79AA1000081	7.88	11.03	9.72	7.09	9.28	7.22	5.68	4.38	3.88	**	-	-
Y79AA1000127	8	5.82	6.2	3.17	4.25	2.35	5.28	8.34	4.47	*	-	-
Y79AA1000130	20.16	11.01	14.68	13.04	8.43	8.05	14.41	16.35	14.16	*	-	-
Y79AA1000131	17.14	8.99	11.18	6.93	10.83	6.92	6.82	8.65	8.52	*	-	-
Y79AA1000134	8.34	7.52	5.68	7.88	5.61	3.09	4.16	2.2	3.07	*	-	-
Y79AA1000143	8.89	7.38	8.14	3.93	5.83	3.75	3.23	2.53	3.96	*	-	-
Y79AA1000144	11.78	8.62	9.18	5.64	6.37	5.58	3.97	3.84	3.38	*	-	-
Y79AA1000150	28.57	28.05	20.04	12.4	17.77	10.86	24.46	18.09	20.29	*	-	-
Y79AA1000153	11.32	11.62	7.41	5.89	9.38	6.53	8.84	5.59	8.52	*	-	-
Y79AA1000165	13.75	6.3	9.45	4.18	5.68	2.88	7.42	8.85	8.78	*	-	-
Y79AA1000175	13.15	8.67	7.74	7.91	7.5	4.67	12.52	5.81	9.18	*	-	-
Y79AA1000181	14.37	9.22	8.77	6.64	8.41	3.83	3.41	3.97	4.85	*	-	-
Y79AA1000202	22.9	17.78	19.17	14.55	12.31	8.38	18.27	13.59	20.94	*	-	-
Y79AA1000207	52.28	63.3	44.9	24.04	38.89	23.79	15.85	13.31	14.27	*	-	-
Y79AA1000214	9.44	4.9	5.68	4.71	4.79	4.18	5.54	2.28	3.2	*	-	-
Y79AA1000222	11.34	8.61	7.85	5.13	7.04	2.28	4.7	3.53	4.67	*	-	-
Y79AA1000225	27.21	25.59	19.24	12.19	15.74	11.35	24.75	16.89	21.48	*	-	-
Y79AA1000227	9.07	8.67	8.62	3.68	4.45	3.9	7.4	6.45	7.51	*	-	-
Y79AA1000230	32.97	25.47	29.26	12.55	10.05	6.88	4.18	3.82	4.93	**	-	-
Y79AA1000231	65.27	41.2	48.87	47.4	47.89	24.84	28.97	32.14	28.29	*	-	-
Y79AA1000239	74.35	85.68	67.35	68.07	52.75	45.41	64.49	54.97	44.91	*	-	-
Y79AA1000258	8.42	5.48	5.2	4.38	4.01	3.09	3.34	2.93	4.62	*	-	-
Y79AA1000268	80.53	76.19	49.38	22.18	30.03	27.42	30.39	28.87	31.5	*	-	-
Y79AA1000269	73.43	71.24	52.77	41.13	70.01	38.67	77.91	62.7	65.45	*	-	-
Y79AA1000270	10.85	7.03	4.1	4.98	7.95	3.4	5.03	5.7	4.72	*	-	-
Y79AA1000280	25.49	25.11	28.88	8.6	12.9	9.27	5.74	8.76	6.26	**	-	-

152 / 157

2743

2744

Y79AA1000285	22.27	18.31	18.83	6.85	10.3	7.36	15.47	22.31	18.9	**	-	-
Y79AA1000286	4.35	2.68	5.01	1.58	8.76	2.11	2.3	2.59	2.22	*	-	-
Y79AA1000307	6.82	4.93	8.12	6.39	9.3	7.38	3	5.88	4.49	*	-	-
Y79AA1000313	4.28	2.34	3.94	1.68	3.1	3.25	1.93	2.47	1.62	*	-	-
Y79AA1000314	1.57	1.83	4.11	1.4	1.41	0.88	0.87	0.14	1.73	*	-	-
Y79AA1000328	3.78	3.39	5.43	2.09	10.7	2.68	3.27	3.33	3.32	*	-	-
Y79AA1000334	9.59	9.05	7.49	5.79	6.64	1.5	5.12	3.62	4.02	* **	-	-
Y79AA1000342	17.48	13.29	18.31	13.02	15.9	13.87	34.2	31.79	30.43	**	-	+
Y79AA1000346	17.08	11.48	18.59	8.72	8.82	6.74	5.35	5.97	4.15	* **	-	-
Y79AA1000347	26.82	26.71	31.25	22.14	31.96	28.74	20.9	20.82	19.69	**	-	-
Y79AA1000349	15.95	20.35	20.7	17.44	19.4	18.58	6.37	7.08	5.15	**	-	-
Y79AA1000355	11.9	11.6	17.89	14.98	13.32	9.11	5.25	6.64	7.85	*	-	-
Y79AA1000368	5.31	4.56	7.88	4.89	6.47	3.77	2.77	3.21	1.85	*	-	-
Y79AA1000388	9.85	11.63	11.71	9.62	9.75	4.51	5.03	4.67	5.04	**	-	-
Y79AA1000392	13.05	13.68	11.25	7.92	12.08	6.34	12.1	13.07	13.52	*	-	-
Y79AA1000405	100.68	82.31	79.28	103.87	109.59	81.35	71.19	52.24	65.44	*	-	-
Y79AA1000410	8.13	6.1	10.22	6.13	5.36	5.28	4.83	4.8	6.17	*	-	-
Y79AA1000420	11.1	11.77	13.48	5.41	8.52	5.43	8.77	7.21	5.64	* **	-	-
Y79AA1000423	7.81	9.21	8.98	4.48	6.29	5.4	4.36	2.75	3.4	* **	-	-
Y79AA1000426	5.75	6.98	10.06	4.68	5.08	3.97	3.38	2.69	2.24	*	-	-
Y79AA1000432	5.62	5.18	10.2	4.71	5.75	4.03	2.85	3.99	8.12	*	-	-
Y79AA1000453	32.42	33.52	37.01	20.21	25.21	18.7	38.98	32.84	33.05	**	-	-
Y79AA1000465	4.43	1.91	3.14	1.7	4.29	1.76	2.57	2.29	2.11	*	-	-
Y79AA1000489	35.31	28.37	27.42	30.17	34.28	32.32	21.89	31.66	24.15	*	-	-
Y79AA1000480	11.78	8.51	13.38	6.22	5.02	6.9	5.84	10.93	8.28	*	-	-
Y79AA1000502	10.24	16.15	21.16	8.68	14	12.35	16.39	18.08	18.17	*	-	-
Y79AA1000521	75.16	71.09	84.73	74.85	78.28	67.85	48.04	51.73	45.89	**	-	-
Y79AA1000534	13.31	15.15	20.4	13.25	14.05	10.71	13.15	11.71	11.75	*	-	-
Y79AA1000538	12.38	14.67	20.45	9.58	11.27	6.96	3.97	0	4.64	**	-	-
Y79AA1000539	15.51	16.32	14.89	9.48	12.74	10.1	5.39	5.2	6.13	* **	-	-
Y79AA1000540	10.2	7.44	12.68	3.41	6.65	4.35	12.15	10.44	10.32	*	-	-
Y79AA1000560	180.34	128.09	149.25	182.08	194.08	150.68	118.75	125.1	108.25	*	-	-
Y79AA1000574	12.15	9.72	10.83	8.68	8.8	4.53	10.59	16.21	10.56	*	-	-
Y79AA1000584	40.28	35.5	37.1	23.89	28.68	25.51	26.95	34.59	37.41	**	-	-
Y79AA1000589	9.65	4.15	9.45	2.79	3.53	2.1	3.54	8.38	4.6	*	-	-
Y79AA1000593	34.28	29.42	37.32	17.43	15.95	16.45	24.99	27.55	21.09	**	-	-
Y79AA1000600	44.58	45.66	51.58	51.4	52.23	39.65	32.05	30.08	26.05	**	-	-
Y79AA1000609	7.91	8.61	8.78	1.95	5.66	3.3	3.74	4.38	7.05	*	-	-
Y79AA1000618	20.27	20.13	21.51	6.81	13.31	6.58	19.85	14.92	16.1	**	-	-
Y79AA1000627	18.35	18.53	18.66	9.08	10.58	9.21	14.78	8.21	19.69	**	-	-
Y79AA1000636	5.21	2.62	7.03	3.38	4.48	4.2	2.54	3.13	1.94	*	-	-
Y79AA1000649	19.51	21.35	25.26	12.58	12.75	9.69	16.89	18.63	18.13	**	-	-
Y79AA1000656	7.37	5.14	7.46	2.86	2.97	2.28	2.83	3.72	2.99	**	-	-
Y79AA1000673	8.38	7.17	11.58	4.74	3.99	5.74	1.87	4.27	2.18	* *	-	-
Y79AA1000674	131.64	157.11	127.73	143.89	140.32	116.2	111.11	118.96	120.05	*	-	-
Y79AA1000678	22.12	27.62	25.85	12.61	21.55	13.08	21.7	19.94	19.35	*	-	-
Y79AA1000682	14.62	11.88	13.44	6.55	8.13	8.38	12.74	13.35	11.29	**	-	-
Y79AA1000683	37.64	24.8	33.28	19.88	18.2	12.73	33.97	32.73	38.17	*	-	-
Y79AA1000697	46.01	35.33	48.41	23.15	27.48	23.15	15.24	20.88	15.95	* **	-	-
Y79AA1000700	3.64	3.69	5.68	3.34	3.08	2.62	3.53	2.27	2.51	*	-	-
Y79AA1000702	60.8	33.01	38	23.71	24.98	18.05	18.21	11.97	15.35	* *	-	-
Y79AA1000704	93.88	71.31	88.5	110.92	93.18	106.88	74.22	61.48	48.95	*	-	-
Y79AA1000705	6.23	6.74	7.37	1.82	2.22	2.09	2.28	0.28	2.39	**	-	-
Y79AA1000717	8.36	6.77	8.13	4.38	8.98	4.55	2.44	3.74	2.84	**	-	-
Y79AA1000722	19.83	11.52	10.81	4.97	5.79	3.88	6.28	5.41	5.45	**	-	-
Y79AA1000724	42.59	35.15	44.52	31.42	30.43	23.04	24.2	29.89	31.58	*	-	-
Y79AA1000726	8.78	4.43	7.31	5.15	4.77	4.25	2.41	3.94	2.09	*	-	-
Y79AA1000734	5.72	4.05	6.45	2.81	2.76	3.48	2.88	2.98	0.88	* *	-	-
Y79AA1000748	7.45	7.72	6.69	2.89	2.82	2.75	2.52	2.6	2.35	**	-	-
Y79AA1000750	41.52	33.1	33.65	18.03	22.85	18.6	28.15	29.63	22.6	**	-	-
Y79AA1000752	9.11	9.26	10.14	4.75	3.47	2.23	2.34	2.01	1.29	**	-	-
Y79AA1000774	19.25	19.31	26.89	10.74	13.71	10.53	12.54	14.59	12.73	*	-	-
Y79AA1000776	12.05	9.97	9.69	6.12	8.81	2.36	10.58	10.24	7.92	*	-	-
Y79AA1000777	17.61	11.59	13.15	7.32	8.15	7.21	6.02	5.12	6.68	*	-	-

153 / 157

2745

2746

Y79AA1000781	20.06	13.02	11.66	10.8	9.69	8.28	18.52	13.51	12.98				
Y79AA1000782	11.43	7.55	9.19	7.76	8.45	7.38	6.82	5.21	4.82				
Y79AA1000784	13.14	10.21	12.84	4.67	5.08	6.71	10.7	9.6	9.71	00	-	-	-
Y79AA1000784	46.19	38.88	38.82	22.35	24.71	16.75	37.28	32.24	34.17	00	-	-	-
Y79AA1000800	7.18	3.91	5.09	0.48	0.36	0.17	2.34	0	0	00	0	-	-
Y79AA1000802	11.45	13.12	10.85	5.78	8.8	4.98	3.89	4.75	3.77	0	00	-	-
Y79AA1000805	21.22	18.33	12.77	8.7	11.14	4.1	15.87	12.52	16.72	0	-	-	-
Y79AA1000814	27.49	21.74	18.65	16.59	17.35	12.98	22	17.22	22.17				
Y79AA1000823	14.24	7.78	9.85	4.94	4.36	5.06	6.79	6.95	8.96	0	-	-	-
Y79AA1000824	16.24	10	9.75	8.11	8.28	6.16	6.82	5.83	8.72				
Y79AA1000827	9.71	3.82	4.8	2.63	3.22	2.56	3.51	2.06	2.83				
Y79AA1000831	7.38	5.79	7.27	4.13	3.32	2.68	4.43	0.57	1.26	00	0	-	-
Y79AA1000833	22.35	19.25	16.88	12.38	7.9	8.89	19.38	11.88	18.04	00	-	-	-
Y79AA1000850	33.44	40.87	26.99	19.52	21.81	14.62	28.93	23.5	32.01	0	-	-	-
Y79AA1000856	6.26	7.4	4.9	2.81	6.29	2.12	4.31	2.45	3.63	0	-	-	-
Y79AA1000882	8.84	4.04	5.06	7.04	4.31	3.07	5.61	5.34	5.05				
Y79AA1000876	21.17	10.25	11.73	7.67	6.82	6.01	5.70	7.01	8				
Y79AA1000888	32.02	25.84	21.68	17.64	17.29	12.59	18.41	20.33	23.17	0	-	-	-
Y79AA1000902	6.26	5.88	6.17	4.27	4.4	2.72	2.73	2.97	2.19	0	00	-	-
Y79AA1000915	76.11	52.08	52.32	70.45	71.68	69.22	39.77	35.2	32.3	0	-	-	-
Y79AA1000959	48.22	40.7	37.08	25.45	33.48	19.11	26.33	21.44	21.93	0	00	-	-
Y79AA1000962	46.82	45.31	38.99	22.79	24.29	17.48	60.82	38.12	53.14	00	-	-	-
Y79AA1000983	25.33	35.17	19.37	13.71	17.82	12.79	14.75	14.39	13.92				
Y79AA1000986	41.31	37.88	31.25	27.6	23.69	23.7	32	18.68	25.22	0	-	-	-
Y79AA1000987	19.13	9.69	10.03	8.88	7.99	4.85	1.27	8.2	0.25				
Y79AA1000988	25.81	18.48	14.41	11.57	12.79	6.94	25.2	19.6	28.68				
Y79AA1000989	40.02	39.87	33.37	23.55	39.74	38.04	32.93	30.52	18.66				
Y79AA1000976	22.75	15.17	16.99	14.98	19.84	15.88	14.85	10.98	11.26				
Y79AA1000975	8.83	7.09	6.8	4.57	5.39	2.65	4.35	3.43	4.33	0	00	-	-
Y79AA1000985	8.02	3.9	5.83	4.32	4.39	1.76	4.37	2.17	1.99				
Y79AA1000989	285.68	243.92	218.77	78.52	138.6	77.87	70.03	48.49	61.43	00	00	-	-
Y79AA1000991	12.26	8.35	10.64	5.63	5.93	8.76	7.23	8.65	6.01				
Y79AA1001013	42.37	37.07	51.31	24.72	27.17	18.23	24.27	34.59	22.31	0	0	-	-
Y79AA1001014	16.53	17.99	19.41	12.93	20.24	14.74	13.35	18.92	15.61				
Y79AA1001019	8.07	6.21	8.29	4.14	6.06	5.04	2.84	5.3	3.61	0	0	-	-
Y79AA1001020	13.89	16.72	13.88	21.03	12.53	10.78	9.5	10.16	11.46	0	-	-	-
Y79AA1001023	7.28	6.24	9.28	4.31	6.03	5.29	3.01	3.87	2.61	0	-	-	-
Y79AA1001030	26.41	43.95	33.88	34.62	21.08	22.74	18.86	22.85	18.21	0	-	-	-
Y79AA1001035	10.6	13.73	10.93	9.68	10.73	7.69	7.26	7.72	8.91	0	-	-	-
Y79AA1001041	14.5	11.14	15.3	8.73	9.88	7.28	9.17	13.51	15.31	0	-	-	-
Y79AA1001043	22.79	19.12	22.36	8.78	13.48	10.29	9.42	13.88	17.78	00	0	-	-
Y79AA1001048	12.86	13.5	12.88	8.85	11.19	7.59	6.44	7.55	8.3	0	00	-	-
Y79AA1001056	6.67	9.77	12.37	5.42	9.6	6.82	2.1	3.99	2.72	0	-	-	-
Y79AA1001061	11.12	17.1	15.18	5.75	10.72	7.78	8.04	10.38	12.77	0	-	-	-
Y79AA1001062	4.7	4.98	6.24	2.95	3.71	2.66	2.82	2.19	2.28	0	00	-	-
Y79AA1001068	6.22	3.69	5.75	4.34	5.88	4.49	3.29	3.38	3.07				
Y79AA1001073	17.6	17.58	20.08	5.98	13.24	7.19	16.17	13.71	20.42	0	-	-	-
Y79AA1001077	7.3	2.88	6.26	7.08	4.43	3.45	1.88	5.22	1.86				
Y79AA1001078	7.29	5.05	6.25	3.73	4.63	2.99	3.3	5.06	3.66	0	-	-	-
Y79AA1001081	7.84	5.5	8.85	3.85	3.81	3.52	2.8	3.1	3.74	0	0	-	-
Y79AA1001088	38.17	40.23	42.34	24.5	22.27	19	24.35	30.94	32.13	00	0	-	-
Y79AA1001089	332.88	368.62	308.34	284.38	330.54	406.65	237.38	378.45	382				
Y79AA1001090	7.05	4.26	8.86	4.12	3.38	2.71	3.39	1.8	2.5				
Y79AA1001105	22.63	22.64	23.57	8.51	14.07	9.94	17.11	15.32	15.62	00	00	-	-
Y79AA1001142	114.5	98.72	131.88	102.55	114.58	143.24	69.56	60.81	74.51	0	-	-	-
Y79AA1001145	98.33	88.67	100.63	125.38	121.38	91.71	78	59.2	74.76	00	-	-	-
Y79AA1001162	126.48	83.55	107.65	92.38	92.62	73.32	87.58	124.59	105.84				
Y79AA1001167	12.61	7.85	13.08	7.92	8.82	6.68	7.15	16.98	9.58				
Y79AA1001176	76.12	80.1	84.48	76.52	85.54	84.34	48.71	59.03	47.45	00	-	-	-
Y79AA1001177	8.46	7.48	10.23	4.08	6.51	3.51	3.45	4.68	2.96	0	-	-	-
Y79AA1001179	64.1	53.61	67.63	28.64	33.09	26.42	13.62	47.33	10.91	00	-	-	-
Y79AA1001185	32.79	30.94	36.04	15.68	21.34	18.32	19.9	17.65	25.06	00	00	-	-
Y79AA1001201	45.08	50.33	41.08	13.71	31.35	17.81	48.18	43.78	38.95	0	-	-	-
Y79AA1001205	18.73	12.88	21.83	8.38	8.48	6.14	7.31	11.85	11.81	0	-	-	-

154 / 157

2747

2748

Y79AA1001211	21.16	21.68	22.83	13.08	12.91	10.43	0.44	20.18	19.17	**	-
Y79AA1001212	128.68	116.33	107.84	161.9	161.84	105.91	80.87	83.57	87.14	**	**
Y79AA1001215	31.68	27.07	33.84	18.3	18.09	14.83	17.05	18.24	23.12	**	*
Y79AA1001228	7.97	5.25	12.95	1.35	3.4	3.64	3.41	10.17	4.89		
Y79AA1001233	8.26	6.53	18.79	1.45	3.77	2.51	2.28	3.79	3.13		
Y79AA1001236	8.84	5.25	11.14	5.02	5.63	3.13	4.08	4.7	2.24		
Y79AA1001239	74.22	59.48	53.54	45.62	83.11	49.05	50.35	42.44	46.18		
Y79AA1001240	70.83	49.07	64.4	30.83	35.04	20.32	48.49	59.48	54.87	*	-
Y79AA1001255	8.89	10.29	10.55	7.47	6.78	6.42	0.72	6.35	0.76	**	*
Y79AA1001264	14.2	10.19	13.36	9.71	10.05	8.87	4.58	4.84	8.39	**	**
Y79AA1001272	16.18	12.6	18.6	1.77	6.76	5.59	8.88	7.59	5.53	**	**
Y79AA1001281	8.87	7.72	6.68	4.17	4.8	3.47	2.65	2.22	2.28	**	**
Y79AA1001299	9.39	10.08	9.8	4.21	5.3	4.49	3.78	3.68	3.29	**	**
Y79AA1001312	40.47	42.72	35.81	17.92	27.92	14.77	29.7	20.85	26.87	*	*
Y79AA1001319	31.92	23	20.23	23.19	28.4	16.88	30.43	20.02	18		
Y79AA1001323	32.99	27.82	34.32	14.07	18.03	14.31	25.9	23.32	27.9	**	-
Y79AA1001328	28.33	16.1	28.18	14.58	12.93	12.88	4.94	7.51	4.42	*	-
Y79AA1001343	12.58	8.01	10.89	6.43	7.59	7.49	6.83	9.08	8.27		
Y79AA1001351	6.64	4.21	8.23	2.99	1.8	1.8	2.08	2.52	1.89	*	*
Y79AA1001364	8.06	6.02	8.85	3.26	2.98	2.12	4.08	2.93	1.74	*	*
Y79AA1001367	145.27	118.42	143.46	117.7	122.8	94.77	76.85	57.81	68.89	**	**
Y79AA1001384	45.86	52.17	41.89	27.43	34.83	21.42	31.33	29.88	26.76	*	-
Y79AA1001381	10.98	12.22	10.84	2.99	9.55	4.77	7.43	7.31	6.01	**	-
Y79AA1001384	24.73	14.05	14.28	9.7	6.32	9.07	8.42	10.16	9.01		
Y79AA1001402	12.42	7.77	9.12	6.82	4.8	4.87	4.74	3.87	4.52	*	*
Y79AA1001410	6.24	2.77	3.09	3.28	1.53	1.95	2.62	0.78	1.99		
Y79AA1001414	27.49	18.53	20.44	10.78	9.47	10.2	16.16	13.43	11.82	*	*
Y79AA1001425	271.77	221.23	225.78	200.18	221.07	217.77	202.63	211.55	197.64		
Y79AA1001427	15.18	10.97	10.9	6.77	10.81	5.39	6.71	6.07	6.74		
Y79AA1001430	22.26	17.58	19.52	10.25	13.63	7.85	11.32	11.76	10.84	*	**
Y79AA1001439	6.78	4.58	4.31	2.6	4.23	1.25	2.94	2.35	2.88	*	-
Y79AA1001485	17.57	9.99	14.92	13.28	10.58	8.25	5.32	10.37	14.36		
Y79AA1001493	20.47	12.23	13.25	8.02	11.27	8.84	13.37	13.83	16.06		
Y79AA1001511	14.52	8.05	9.32	8.33	6.58	5.82	5.2	6	5.98		
Y79AA1001523	10.03	6.72	7.33	8.48	4.81	3.84	4.54	2.21	2.59	*	-
Y79AA1001530	11.44	7.57	8.59	8.84	8.84	5.29	6.2	7.76	5.77		
Y79AA1001532	13.66	14.73	11.84	5.35	5.16	4.93	5.08	4.52	6.49	**	**
Y79AA1001533	68.6	68.51	50.9	29.97	36.82	26.92	59.28	61.19	56.58	*	-
Y79AA1001541	8.12	8.32	5.25	4.02	4.89	2.24	4.69	2.24	2.18	*	-
Y79AA1001548	67.38	47.65	44.48	30.51	28.42	21.82	46.07	51.22	51.86	*	-
Y79AA1001555	14.35	9.76	8.13	8.86	4.86	4.97	8.19	7.68	10.88		
Y79AA1001562	18.44	13.73	12.82	8.99	10.8	7.19	10.2	8.39	15.75	*	-
Y79AA1001581	24.34	18.91	17.82	9.8	12.04	10.4	15.8	9.98	16.33	*	-
Y79AA1001585	30.32	20.83	22.43	20.22	25.12	15.45	0.89	15.13	0.14	*	-
Y79AA1001592	28.53	25.65	26.36	28.33	18.68	7.28	20.69	8.44	14.72	*	-
Y79AA1001594	23.28	21.85	24.28	14.25	15.59	10.11	22.55	14.53	20.27	**	-
Y79AA1001603	12.47	11.04	10.84	8.8	11.48	6.78	22.63	14.12	27.5		
Y79AA1001612	25.09	18.27	18.81	23.47	18.84	13.34	20.3	16.35	18.99		
Y79AA1001630	22.84	13.91	12.22	13.14	13.53	8.02	11.73	12.98	17.87		
Y79AA1001647	33.54	24.86	22.88	23.22	25.84	18.49	26.93	21.3	26.55		
Y79AA1001684	12.2	10.58	13.13	7.78	6.05	5.54	10.34	8.1	11.69	**	-
Y79AA1001685	11.52	9	9.75	8.87	7.31	5.13	5.82	6.12	7.89	*	*
Y79AA1001679	24.51	18.98	17.88	11.13	12.84	9.27	24.02	13.28	18.8	*	-
Y79AA1001692	7.23	6.72	8.1	2.24	2.83	2.63	3.64	2.31	3.08	**	**
Y79AA1001696	14.23	12.31	9.21	5.29	8.48	4.81	14.27	13.28	13.01	*	-
Y79AA1001705	12.04	11.4	8.55	9.78	11.8	6.21	4.61	5.98	4.57	**	-
Y79AA1001711	13.72	7.72	10.63	6.49	4.11	3.08	6.08	4.08	7.4	*	-
Y79AA1001717	4.71	1.65	3.26	3.34	1.71	0.7	1.49	1.63	1.39		
Y79AA1001719	22.12	12.72	16.33	9.89	12.72	6.53	12.21	11.1	11.78		
Y79AA1001727	38.98	27.16	22.35	13.65	17.35	9.31	38.01	28.39	34.23	*	-
Y79AA1001750	21.5	17.52	18.42	10.15	15.4	8.01	11.94	9.9	11.41	*	**
Y79AA1001750	127.24	115.03	77.78	58.28	73.8	43.62	140.25	100.84	138.73	*	-
Y79AA1001777	8.77	8.32	6.97	2.57	5.24	2.46	5.26	4.15	4.71	*	**
Y79AA1001781	2.86	2.24	2.15	1.61	1.88	2.55	0.74	1.53	1.3	*	-

155 / 157

【0979】

【表508】

2750

156 / 157

7751

Y79AA1002208	7.86	6.79	4.88	2.41	3.6	5.74	5.49	1.51	2.0				
Y79AA1002208	21.01	17.64	15.14	5.6	4.07	5.57	8.81	5.17	2.65	**	**	-	-
Y79AA1002209	14.82	11.28	11.88	8.23	4.7	2.82	4.71	1.33	1.18	**	**	-	-
Y79AA1002210	13.64	7.39	7.59	9.08	4.82	5.18	20.5	2.05	6.37				
Y79AA1002211	11.78	10.59	13.47	10.43	8.65	6.52	12.32	8.42	11.25				
Y79AA1002213	40.78	31.99	22.86	18.41	26.57	14.98	45.88	32.4	41.97				
Y79AA1002215	64.92	41.69	39.55	24.88	24.38	11.28	37.49	23.8	35.98	*			
Y79AA1002220	17.03	11.5	20.58	7.13	6.65	5.31	4.57	4.8	5.51	*	*	-	-
Y79AA1002226	48.65	31.27	31.34	7.35	12.72	13.65	9.19	6.65	11.13	*	**	-	-
Y79AA1002229	7.88	8.84	5.37	6.02	4.67	2.85	3.67	2.52	3.73	*			
Y79AA1002234	20.83	13.27	12.39	9.34	6.36	3.6	6.9	3.38	5.84	*	*	-	-
Y79AA1002235	28.03	23.84	21.24	15.07	14.67	9.39	10.75	8.42	13.64	*	**	-	-
Y79AA1002246	9.72	14.9	10.35	5.25	6.99	4.12	8.31	3.82	6.69	*			
Y79AA1002258	12.35	12.02	7.88	7.82	11.67	8.56	9.77	4.5	4.81				
Y79AA1002279	51.56	40.19	41.11	6.28	2.78	2.07	15.99	20.03	22.13	**	**	-	-
Y79AA1002292	13.64	7.58	4.14	5.71	4.43	2.94	6.45	8.01	8.38				
Y79AA1002288	9.43	8.29	4.77	3.29	5.48	4.2	4.42	4.85	3.79				
Y79AA1002307	9.31	8.07	6.82	3.9	4.4	2.44	5.17	1.78	2.58	*	*	-	-
Y79AA1002309	8.88	7.96	8.79	3.32	4.28	2.83	4.23	2.88	3.97	**	**	-	-
Y79AA1002311	16.61	9.27	11.76	6.8	6.2	3.45	7.45	3.09	6.4				
Y79AA1002334	13.06	8.36	7.7	5.13	4.89	3.33	5.91	4.32	5.92	*			
Y79AA1002351	13.81	12.49	9.42	7.1	1.6	6.19	7.95	4.93	5.8	*	*	-	-
Y79AA1002355	31.74	30.6	21.85	12.21	15.81	9.74	20.54	18.29	18.48	*	*	-	-
Y79AA1002361	23.42	15.4	18.02	12.63	10.73	6.85	25.86	17.1	25.6	*			
Y79AA1002365	12.42	6.37	7.19	3.15	4.11	3.03	4.29	4.74	4.01				
Y79AA1002373	8.95	6.89	5.48	5.13	4.81	3.4	9.6	4.57	7.84				
Y79AA1002376	1550.5	2589.2	1850.6	462.62	827.86	616.71	1477.8	1040.9	1062.1	*			
Y79AA1002378	20.24	17.32	13.34	5.14	9.41	4.23	19.28	11.07	16.88	*			
Y79AA1002381	116.11	128.86	74.48	110.68	141.78	92.68	155.95	123.08	170.94				
Y79AA1002388	32.4	32.3	27.31	13.85	28.76	11.62	21.29	16.32	21.24	*			
Y79AA1002399	11.13	8.22	7.72	4.26	5.54	4.87	7.58	6.9	8.25	*			
Y79AA1002407	12.68	14.43	18.13	7.72	14.18	6.84	5.83	9.78	4.59	*			
Y79AA1002413	16.98	12.77	14.95	6.14	9.13	4.52	8.44	10.73	7.99	*	*	-	-
Y79AA1002416	7.52	8.19	8.76	5.47	10.72	5.8	8.2	8.05	6.59	*			
Y79AA1002429	17.73	18.81	8.81	5.82	10.24	4.73	3.65	8.89	5.68	*			
Y79AA1002431	3.38	3.05	6.2	3.01	5.89	1.9	2.81	2.79	1.69				
Y79AA1002433	0.94	11.67	9.29	5.11	5.57	3.18	3.49	4.8	3.87	**	**	-	-
Y79AA1002445	33.47	25.62	23.49	15.99	10.87	7.02	18.92	25.26	13.87	*			
Y79AA1002461	7.94	6.22	7.84	3.35	7.35	4.7	3.49	2.25	3.65	**			
Y79AA1002480	778.44	339.4	681.02	542.58	498.15	389	592.67	971	768.71				
Y79AA1002471	11.38	8.13	15.35	12.81	13.4	11.43	4.84	8.06	4.47	*			
Y79AA1002472	31.22	33.08	31.17	18.15	21.85	9.34	19.29	20.14	20.03	*	**	-	-
Y79AA1002474	30.68	12.29	10.71	6.77	7.3	7.75	3.17	7.37	4.88	*	*	-	-
Y79AA1002482	10.09	53.68	38.63	19.82	23.45	17.38	21.9	25.81	23.08	**	*	-	-
Y79AA1002487	8.33	8.29	7.43	7.28	8.45	6.44	5.34	3.78	3.88	*			
Y79AA1002493	143.18	108.89	117.63	56.22	71.49	57.31	59.78	51.39	62.37	**	**	-	-
Y79AA1002493	54.75	41.58	40.36	30.64	28.52	19.33	38.02	48.19	48.7	**	*	-	-
ZR9GC1006276	4.26	7	6.52	3.16	2.97	2.19	2.99	2.5	2.72	**	**	-	-

7752

【0981】実施例16. オリゴキャップ法で作製されたcDNAライブラリーからの新規cDNAクローンの選抜
オリゴキャップ法で作製された cDNAライブラリーから得られたクローン の5'末端配列中に開始コドンから始まる50アミノ酸以上の蛋白質コード領域が推定される54クローン、HEMBA1000497、HEMBA1001750、HEMBA1003854、HEMBA1004193、HEMBA1004860、HEMBA1005572、HEMBA1006038、HEMBA1006092、HEMBA1006406、HEMBA1006650、HEMBA1006812、HEMBA100672、HEMBA1001197、HEMBA1001871、HEMBA1001252、HEMBA1002094、NT2RM4000634、NT2RM4000657、NT2RM4000783、NT2RM4000857、NT2RM4001178、NT2RM4002420、NT2RP2000198、NT2RP2000551、NT2RP2000660、NT2RP2001214、NT2RP2001460、NT2RP2001756、NT2RP2002056、NT2RP2002677、NT2RP2002755、NT2RP2002843、NT2RP2003101、NT2RP2003799、NT2RP2004095、NT2RP2004732、NT2RP2004920、NT2RP2005454、NT2RP2005776、NT2RP2005806、NT2RP2005882、NT2RP3001282、NT2RP3001723、NT2RP3002099、NT2RP3003155、NT2RP3004028、OVARC1000008、OVARC1000724、OVARC1000751、OVARC1001029、PLACE1000814、PLACE1003030、PLACE1005549、PLACE1007218を新たに選抜した。中でもHEMBA1000497、HEMBA1003854、HEMBA1004193、HEMBA1006812、HEMBA1001871、NT2RM4000657、NT2RM4001178、NT2RP2001756、NT2RP2002677、NT2RP2002755、NT2RP

2002843、NT2RP2004095、NT2RP2004920、NT2RP200580
6、NT2RP3001282、NT2RP3002099、NT2RP3003155、OVARC
1000724、OVARC1001029、PLACE1000814、PLACE100303
0、PLACE1005549、PLACE1007218の23クローンは100アミ
ノ酸以上の蛋白質コード領域が推定され、蛋白質コード
領域を含む確率が非常に高いことが示された。

【0982】各クローンの最大ATGpr1値を表510に示す。HEMBA1006812、HEMBA1001871、NT2RP3001282は最大ATGpr1値が0.3より大であるため全長である確率は高いが、その他HEMBA1000497、HEMBA1001750、HEMBA1003854、HEMBA1004193、HEMBA1004860、HEMBA1005572、HEMBA1006038、HEMBA1006092、HEMBA1006406、HEMBA1006650、HEMBA1006672、HEMBA1001197、HEMBA1001252、HEMBA1002094、NT2RP4000634、NT2RP4000657、NT2RP4000783、NT2RP4000857、NT2RP4001178、NT2RP4002420、NT2RP2000198、NT2RP2000551、NT2RP2000660、NT2RP2001214、NT2RP2001460、NT2RP2001756、NT2RP2002056、NT2RP2002677、NT2RP2002755、NT2RP2002843、NT2RP2003101、NT2RP2003799、NT2RP2004095、NT2RP2004732、NT2RP2004920、NT2RP2005454、NT2RP2005776、NT2RP2005806、NT2RP2005882、NT2RP3001723、NT2RP3002099、NT2RP3003155、NT2RP3004028、OVARC1000008、OVARC1000724、OVARC1000751、OVARC1001029、PLACE1000814、PLACE1003030、PLACE1005549、PLACE1007218は最大ATGpr1値

2753

が0.3以下であるため全長率が低いとされるものの依然全長である可能性はある。表511(表2と同じ内容を表す)中に54クローンにおける5'側と3'側を規定する塩基配列を示す配列番号と、そのポリヌクレオチドをインサートとして含む実施例で取得したプラスミドクロンの対応関係を示す。5'配列 配列名の右側が5'配列の配列番号であり、3'配列 配列名の右側が3'配列の配列番号である。選抜された54クローンの5'末端配列をSwiss-Protに対して、5'末端配列、3'末端配列をGenBank、UniGeneに対して相溶性検索した。それぞれの検索結果は、明細書の最後に相溶性検索結果データ7~11として示した。

【0983】得られたトップヒットデータからは、少なくとも、分泌・膜蛋白質、糖蛋白質、シグナル伝達関連蛋白質、転写関連蛋白質、疾患関連蛋白質、蛋白質合成・輸送関連蛋白質のいずれかのカテゴリーに属する蛋白質をコードすると推定されるクローンが7クローンあった。いずれも該カテゴリーに属する既知蛋白質と比較的相溶性の低いとされるクローンであった。ここでいう「比較的相溶性が低い」とは、「比較的相溶性が高い」条件(Swiss-Protデータベースの既知配列との比較において、Identityが60%以上、かつP値が 10^{-10} 以下のスコアである)は満たさないが、Swiss-Protデータベースの既知配列との比較において、比較配列の長さが55塩基以上、かつIdentityが25%以上、かつP値が 10^{-4} 以下のスコアであることを示す。7クローンのうち、分泌・膜蛋白質のカテゴリーに属する蛋白質をコードすると推定されるクローンはHEMBA1001750、NT2RM4000857の2クローンであった(他のカテゴリーに重複して属するクローン)。糖蛋白質のカテゴリーに属する蛋白質をコードすると推定されるクローンはHEMBA1001750、NT2RM4000857の2クローンであった(他のカテゴリーに重複して属するクローン)。シグナル伝達関連蛋白質のカテゴリーに属する蛋白質をコードすると推定されるクローンはPLACE1005549の1クローンであった。転写関連蛋白質のカテゴリーに属する蛋白質をコードすると推定されるクローンはHEMBA1005572、NT2RP2001756、NT2RP2005776の3クローンであった。疾患関連蛋白質のカテゴリーに属する蛋白質をコードすると推定されるクローンはNT2RM4000857の1クローンであった(他のカテゴリーに重複して属するクローン)。蛋白質合成・輸送関連蛋白質のカテゴリーに属する蛋白質をコードすると推定されるクローンはHEMBA1001750の1クローンであった(実施例12.参照)。

【0984】

【表510】

実施例16で選抜されたクローンの最大ATGpr1値

クローン名 配列名 最大ATGpr1値

50

2754

HEMBA1000497	F-HEMBA1000497	0.25
HEMBA1001750	F-HEMBA1001750	0.08
HEMBA1003854	F-HEMBA1003854	0.23
HEMBA1004193	F-HEMBA1004193	0.22
HEMBA1004860	F-HEMBA1004860	0.29
HEMBA1005572	F-HEMBA1005572	0.24
HEMBA1006038	F-HEMBA1006038	0.29
HEMBA1006092	F-HEMBA1006092	0.28
HEMBA1006406	F-HEMBA1006406	0.26
HEMBA1006650	F-HEMBA1006650	0.22
HEMBA1006812	F-HEMBA1006812	0.71
HEMBA100672	F-HEMBA100672	0.24
HEMBA1001197	F-HEMBA1001197	0.22
HEMBA1001871	F-HEMBA1001871	0.94
MAMMA1001252	F-MAMMA1001252	0.29
MAMMA1002094	F-MAMMA1002094	0.28
NT2RM4000634	F-NT2RM4000634	0.07
NT2RM4000657	F-NT2RM4000657	0.24
NT2RM4000783	F-NT2RM4000783	0.22
NT2RM4000857	F-NT2RM4000857	0.12
NT2RM4001178	F-NT2RM4001178	0.27
NT2RM4002420	F-NT2RM4002420	0.06
NT2RP2000198	F-NT2RP2000198	0.15
NT2RP2000551	F-NT2RP2000551	0.07
NT2RP2000660	F-NT2RP2000660	0.22
NT2RP2001214	F-NT2RP2001214	0.26
NT2RP2001460	F-NT2RP2001460	0.07
NT2RP2001756	F-NT2RP2001756	0.17
NT2RP2002056	F-NT2RP2002056	0.12
NT2RP2002677	F-NT2RP2002677	0.14
NT2RP2002755	F-NT2RP2002755	0.12
NT2RP2002843	F-NT2RP2002843	0.11
NT2RP2003101	F-NT2RP2003101	0.13
NT2RP2003799	F-NT2RP2003799	0.24
NT2RP2004095	F-NT2RP2004095	0.16
NT2RP2004732	F-NT2RP2004732	0.18
NT2RP2004920	F-NT2RP2004920	0.15
NT2RP2005454	F-NT2RP2005454	0.09
NT2RP2005776	F-NT2RP2005776	0.19
NT2RP2005806	F-NT2RP2005806	0.27
NT2RP2005882	F-NT2RP2005882	0.11
NT2RP3001282	F-NT2RP3001282	0.39
NT2RP3001723	F-NT2RP3001723	0.22
NT2RP3002099	F-NT2RP3002099	0.20
NT2RP3003155	F-NT2RP3003155	0.29
NT2RP3004028	F-NT2RP3004028	0.13
OVARC1000008	F-OVARC1000008	0.23
OVARC1000724	F-OVARC1000724	0.27
OVARC1000751	F-OVARC1000751	0.28
OVARC1001029	F-OVARC1001029	0.25

(1379)

特開2002-191363

2755

2756

PLACE1000814	F-PLACE1000814	0.21	*
PLACE1003030	F-PLACE1003030	0.26	【0985】
PLACE1005549	F-PLACE1005549	0.16	【表511】
PLACE1007218	F-PLACE1007218	0.30	*

クローン名	5'末端配列 配列名	5'末端 配列番号	3'末端配列 配列名	3'末端 配列番号
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HEMBA1000497	F-HEMBA1000497	16111	R-HEMBA1000497	16165
HEMBA1001750	F-HEMBA1001750	16112	R-HEMBA1001750	16166
HEMBA1003854	F-HEMBA1003854	16113	R-HEMBA1003854	16167
HEMBA1004193	F-HEMBA1004193	16114	R-HEMBA1004193	16168
HEMBA1004860	F-HEMBA1004860	16115	R-HEMBA1004860	16169
HEMBA1005572	F-HEMBA1005572	16116	R-HEMBA1005572	16170
HEMBA1006038	F-HEMBA1006038	16117	R-HEMBA1006038	16171
HEMBA1006092	F-HEMBA1006092	16118	R-HEMBA1006092	16172
HEMBA1006406	F-HEMBA1006406	16119	R-HEMBA1006406	16173
HEMBA1006650	F-HEMBA1006650	16120	R-HEMBA1006650	16174
HEMBA1006812	F-HEMBA1006812	16121	R-HEMBA1006812	16175
HEMBB1000672	F-HEMBB1000672	16122	R-HEMBB1000672	16176
HEMBB1001197	F-HEMBB1001197	16123	R-HEMBB1001197	16177
HEMBB1001871	F-HEMBB1001871	16124	R-HEMBB1001871	16178
MAMMA1001252	F-MAMMA1001252	16125	R-MAMMA1001252	16179
MAMMA1002094	F-MAMMA1002094	16126	R-MAMMA1002094	16180
NT2RM4000634	F-NT2RM4000634	16127	R-NT2RM4000634	16181
NT2RM4000657	F-NT2RM4000657	16128	R-NT2RM4000657	16182
NT2RM4000783	F-NT2RM4000783	16129	R-NT2RM4000783	16183
NT2RM4000857	F-NT2RM4000857	16130	R-NT2RM4000857	16184
NT2RM4001178	F-NT2RM4001178	16131	R-NT2RM4001178	16185
NT2RM4002420	F-NT2RM4002420	16132	R-NT2RM4002420	16186
NT2RP2000198	F-NT2RP2000198	16133	R-NT2RP2000198	16187
NT2RP2000551	F-NT2RP2000551	16134	R-NT2RP2000551	16188
NT2RP2000660	F-NT2RP2000660	16135	R-NT2RP2000660	16189
NT2RP2001214	F-NT2RP2001214	16136	R-NT2RP2001214	16190
NT2RP2001460	F-NT2RP2001460	16137	R-NT2RP2001460	16191
NT2RP2001756	F-NT2RP2001756	16138	R-NT2RP2001756	16192
NT2RP2002056	F-NT2RP2002056	16139	R-NT2RP2002056	16193
NT2RP2002677	F-NT2RP2002677	16140	R-NT2RP2002677	16194
NT2RP2002755	F-NT2RP2002755	16141	R-NT2RP2002755	16195
NT2RP2002843	F-NT2RP2002843	16142	R-NT2RP2002843	16196
NT2RP2003101	F-NT2RP2003101	16143	R-NT2RP2003101	16197
NT2RP2003799	F-NT2RP2003799	16144	R-NT2RP2003799	16198
NT2RP2004095	F-NT2RP2004095	16145	R-NT2RP2004095	16199
NT2RP2004732	F-NT2RP2004732	16146	R-NT2RP2004732	16200
NT2RP2004920	F-NT2RP2004920	16147	R-NT2RP2004920	16201
NT2RP2005454	F-NT2RP2005454	16148	R-NT2RP2005454	16202
NT2RP2005776	F-NT2RP2005776	16149	R-NT2RP2005776	16203
NT2RP2005806	F-NT2RP2005806	16150	R-NT2RP2005806	16204
NT2RP2005882	F-NT2RP2005882	16151	R-NT2RP2005882	16205
NT2RP3001282	F-NT2RP3001282	16152	R-NT2RP3001282	16206

2757

NT2RP3001723	F-NT2RP3001723	16153	R-NT2RP3001723	16207
NT2RP3002099	F-NT2RP3002099	16154	R-NT2RP3002099	16208
NT2RP3003155	F-NT2RP3003155	16155	R-NT2RP3003155	16209
NT2RP3004028	F-NT2RP3004028	16156	R-NT2RP3004028	16210
OVARC1000008	F-OVARC1000008	16157	R-OVARC1000008	16211
OVARC1000724	F-OVARC1000724	16158	R-OVARC1000724	16212
OVARC1000751	F-OVARC1000751	16159	R-OVARC1000751	16213
OVARC1001029	F-OVARC1001029	16160	R-OVARC1001029	16214
PLACE1000814	F-PLACE1000814	16161	R-PLACE1000814	16215
PLACE1003030	F-PLACE1003030	16162	R-PLACE1003030	16216
PLACE1005549	F-PLACE1005549	16163	R-PLACE1005549	16217
PLACE1007218	F-PLACE1007218	16164	R-PLACE1007218	16218

2758

【0986】実施例17. 推定アミノ酸配列に対するシグナル配列、膜貫通領域および機能ドメインの検索
全長塩基配列から推定されたアミノ酸配列に対して、アミノ末端のシグナル配列の有無と膜貫通領域の有無を予測、さらに蛋白質の機能ドメイン（モチーフ）検索を行った。アミノ末端のシグナル配列についてはPSORT [K. Nakai & M. Kanehisa, Genomics, 14: 897-911 (1992)] を、膜貫通領域についてはSOSUI [T. Hirokawa et al. Bioinformatics, 14: 378-379 (1998)]（三井情報開発株式会社販売）を用いて解析を行った。機能ドメインの検索についてはPfam (<http://www.sanger.ac.uk/Software/Pfam/index.shtml>) を用いた。PSORTやSOSUIにより、アミノ末端のシグナル配列や膜貫通領域が予測されたアミノ酸配列は分泌、膜蛋白質であると予測された。また、Pfamによる機能ドメイン検索において、ある機能ドメインにヒットしたアミノ酸配列はヒットデータをもとに、例えばPROSITE(<http://www.expasy.ch/cgi-bin/prosite-list.pl>)にある機能カテゴリー分類を参照にしてその蛋白質の機能予測することができる。また、PROSITEでの機能ドメインの検索も可能である。各ソフトウェアによる検索結果を以下に示す。

【0987】PSORTにより推定アミノ酸配列にシグナル配列を検出されたクローンは、以下の通りであった。
HEMBA1001052, HEMBA1001407, HEMBA1002486, HEMBA1002661, HEMBA1002818, HEMBA1002876, HEMBA1003086, HEMBA1003711, HEMBA1004752, HEMBA1005991, HEMBA1006067, HEMBA1006173, HEMBA1006198, HEMBA1006789, HEMBA1006921, HEMBB1000054, HEMBB1000175, HEMBB1002692, MAMMA1000798, MAMMA1002427, MAMMA1002881, MAMMA1003035, NT2RM1000035, NT2RM1000742, NT2RM1000811, NT2RM1000905, NT2RM1001008, NT2RM2000287, NT2RM2000609, NT2RM2001613, NT2RM4000634, NT2RM4000778, NT2RM4002339, NT2RM4002460, NT2RP1000782, NT2RP1000856, NT2RP1001247, NT2RP1001546, NT2RP1001569, NT2RP2001597, NT2RP2002537, NT2RP2004142, NT2RP2005752, NT2RP2005812, NT2RP3001084, NT2RP3001589, NT2RP3002163, NT2RP3002650, NT2RP3003145, NT2RP3003242, 50

NT2RP3003621, NT2RP3004282, NT2RP3004503, NT2RP4000051, NT2RP4000151, NT2RP4000243, NT2RP4000259, NT2RP4000323, NT2RP4000417, NT2RP4001064, NT2RP400117, NT2RP4001730, NT2RP4001739, NT2RP4002075, NT2RP5003500, OVARC1001154, PLACE1000611, PLACE1003030, PLACE1003044, PLACE1003369, PLACE1003596, PLACE1004258, PLACE1005086, PLACE1006239, PLACE1006754, PLACE1006829, PLACE1007954, PLACE1008424, PLACE1008533, PLACE1008693, PLACE1010622, PLACE1010942, PLACE2000176, PLACE2000341, PLACE2000379, PLACE2000427, PLACE2000477, PLACE4000431, PLACE4000593, THYR01000156, THYR01001134, THYR01001287, Y79AA1000258, Y79AA1001874, Y79AA1002399
HEMBA1001871, HEMBB1001925, MAMMA1000778, MAMMA1000897, MAMMA1001080, NT2RP2004300, NT2RP3002985, NT2RP3003059, OVARC1000689, OVARC1000890, PLACE1005162, PLACE3000399, PLACE3000455, PLACE4000247, PLACE4000259, PLACE4000494

【0988】SOSUIにより推定アミノ酸配列に膜貫通領域を検出されたクローンは、以下の通りであった。
HEMBA1000005, HEMBA1000356, HEMBA1000518, HEMBA1000531, HEMBA1000637, HEMBA1000719, HEMBA1000817, HEMBA1000822, HEMBA1000870, HEMBA1000991, HEMBA1001052, HEMBA1001085, HEMBA1001286, HEMBA1001351, HEMBA1001407, HEMBA1001446, HEMBA1001510, HEMBA1001515, HEMBA1001557, HEMBA1001746, HEMBA1002092, HEMBA1002125, HEMBA1002150, HEMBA1002166, HEMBA1002462, HEMBA1002477, HEMBA1002486, HEMBA1002609, HEMBA1002659, HEMBA1002661, HEMBA1002780, HEMBA1002818, HEMBA1002876, HEMBA1002921, HEMBA1003077, HEMBA1003079, HEMBA1003086, HEMBA1003096, HEMBA1003281, HEMBA1003286, HEMBA1003711, HEMBA1003742, HEMBA1003803, HEMBA1004143, HEMBA1004146, HEMBA1004341, HEMBA1004461, HEMBA1004577, HEMBA1004637, HEMBA1004752, HEMBA1004756, HEMBA1004850, HEMBA1004889, HEMBA1004923, HEMBA1004930, HEMBA1005029, HEMBA1005035, HEMBA1005050, HEMBA1005552, HEMBA1005588, HEMBA1005

2759

616, HEMBA1005991, HEMBA1006036, HEMBA1006067, HEMBA1006293, HEMBA1006492, HEMBA1006502, HEMBA1006659, HEMBA1006758, HEMBA1006789, HEMBA1006921, HEMBA1006926, HEMBA1007203, HEMBB1000050, HEMBB1000054, HEMBB1000556, HEMBB1000593, HEMBB1000631, HEMBB1000763, HEMBB1000827, HEMBB1000915, HEMBB1000975, HEMBB1001112, HEMBB1001177, HEMBB1001302, HEMBB1001348, HEMBB1001962, HEMBB1002142, HEMBB1002190, HEMBB1002247, HEMBB1002387, HEMBB1002550, HEMBB1002600, HEMBB1002692, MAMMA1000129, MAMMA1000133, MAMMA1000277, MAMMA1000278, MAMMA1000410, MAMMA1000416, MAMMA1000472, MAMMA1000714, MAMMA1000731, MAMMA1000734, MAMMA1000798, MAMMA1000842, MAMMA1000956, MAMMA1001008, MAMMA1001030, MAMMA1001139, MAMMA1001154, MAMMA1001388, MAMMA1001411, MAMMA1001487, MAMMA1001751, MAMMA1001771, MAMMA1002461, MAMMA1002524, MAMMA1002598, MAMMA1002684, MAMMA1002769, MAMMA1002890, MAMMA1002938, MAMMA1003146, NT2RM1000035, NT2RM1000037, NT2RM1000062, NT2RM1000131, NT2RM1000257, NT2RM1000260, NT2RM1000355, NT2RM1000648, NT2RM1000742, NT2RM1000800, NT2RM1000811, NT2RM1000857, NT2RM1000867, NT2RM1000882, NT2RM1001008, NT2RM1001115, NT2RM1001139, NT2RM2000259, NT2RM2000395, NT2RM2000402, NT2RM2000407, NT2RM2000422, NT2RM2000566, NT2RM2000581, NT2RM2000609, NT2RM2001370, NT2RM2001393, NT2RM2001499, NT2RM2001613, NT2RM2001648, NT2RM2001659, NT2RM2001671, NT2RM2001718, NT2RM2001760, NT2RM2001785, NT2RM2001823, NT2RM2001930, NT2RM2001950, NT2RM2001998, NT2RM2002049, NT2RM4000046, NT2RM4000233, NT2RM4000433, NT2RM4000520, NT2RM4000634, NT2RM4000674, NT2RM4000700, NT2RM4000764, NT2RM4000795, NT2RM4000820, NT2RM4000857, NT2RM4001032, NT2RM4001054, NT2RM4001455, NT2RM4001813, NT2RM4001930, NT2RM4001987, NT2RM4002054, NT2RM4002073, NT2RM4002145, NT2RM4002146, NT2RM4002194, NT2RM4002339, NT2RM4002438, NT2RM4002446, NT2RM4002452, NT2RM4002460, NT2RM4002493, NT2RM4002571, NT2RP1000191, NT2RP1000358, NT2RP1000418, NT2RP1000547, NT2RP1000609, NT2RP1000677, NT2RP1000767, 【0989】NT2RP1000782, NT2RP1000856, NT2RP1001113, NT2RP1001247, NT2RP1001286, NT2RP1001310, NT2RP1001311, NT2RP1001313, NT2RP1001385, NT2RP1001449, NT2RP1001546, NT2RP1001569, NT2RP2000032, NT2RP2000040, NT2RP2000070, NT2RP2000091, NT2RP2000114, NT2RP2000120, NT2RP2000173, NT2RP2000175, NT2RP2000195, NT2RP2000248, NT2RP2000270, NT2RP2000283, NT2RP2000289, NT2RP2000459, NT2RP2000516, NT2RP2000842, NT2RP2000892, NT2RP2001081, NT2RP2001268, NT2RP2001295, NT2RP2001366, NT2RP2001576, NT2RP2001581, NT2RP2001597, NT2RP2001947, NT2RP2001991, NT2RP2

2760

002025, NT2RP2002312, NT2RP2002385, NT2RP2002479, NT2RP2002537, NT2RP2002643, NT2RP2002701, NT2RP2002740, NT2RP2002857, NT2RP2003125, NT2RP2003297, NT2RP2003433, NT2RP2003446, NT2RP2003466, NT2RP2003629, NT2RP2003777, NT2RP2003781, NT2RP2004041, NT2RP2004194, NT2RP2004270, NT2RP2004681, NT2RP2004775, NT2RP2004799, NT2RP2004936, NT2RP2005012, NT2RP2005159, NT2RP2005227, NT2RP2005270, NT2RP2005344, NT2RP2005509, NT2RP2005752, NT2RP2005781, NT2RP2005784, NT2RP2005812, NT2RP2006069, NT2RP2006100, NT2RP2006141, NT2RP2006261, NT2RP2006571, NT2RP3000092, NT2RP3000134, NT2RP3000333, NT2RP3000393, NT2RP3000439, NT2RP3000441, NT2RP3000531, NT2RP3000685, NT2RP3000826, NT2RP3000852, NT2RP3001126, NT2RP3001176, NT2RP3001260, NT2RP3001355, NT2RP3001383, NT2RP3001426, NT2RP3001453, NT2RP3001497, NT2RP3001538, NT2RP3001716, NT2RP3001727, NT2RP3001739, NT2RP3001799, 【0990】NT2RP3001943, NT2RP3001944, NT2RP3002002, NT2RP3002014, NT2RP3002054, NT2RP3002108, NT2RP3002163, NT2RP3002351, NT2RP3002455, NT2RP3002549, NT2RP3002628, NT2RP3002650, NT2RP3002687, NT2RP3002701, NT2RP3002869, NT2RP3002969, NT2RP3003008, NT2RP3003071, NT2RP3003101, NT2RP3003145, NT2RP3003302, NT2RP3003353, NT2RP3003409, NT2RP3003716, NT2RP3003918, NT2RP3004207, NT2RP3004454, NT2RP3004503, NT2RP4000051, NT2RP4000151, NT2RP4000243, NT2RP4000259, NT2RP4000323, NT2RP4000500, NT2RP4000560, NT2RP4000588, NT2RP4000713, NT2RP4000724, NT2RP4000833, NT2RP4000878, NT2RP4000907, NT2RP4000925, NT2RP4000928, NT2RP4000973, NT2RP4000989, NT2RP4001057, NT2RP4001064, NT2RP4001079, NT2RP4001117, NT2RP4001138, NT2RP4001150, NT2RP4001174, NT2RP4001274, NT2RP4001345, NT2RP4001372, NT2RP4001373, NT2RP4001379, NT2RP4001498, NT2RP4001547, NT2RP4001571, NT2RP4001644, NT2RP4001677, NT2RP4001803, NT2RP4001822, NT2RP4001975, NT2RP4002052, NT2RP4002075, NT2RP5003500, NT2RP5003506, NT2RP5003522, NT2RP5003534, OVARC1000151, OVARC1000241, OVARC1000335, OVARC1000700, OVARC1000722, OVARC1000751, OVARC1000850, OVARC1000924, OVARC1000936, OVARC1000959, OVARC1000984, OVARC1001034, OVARC1001129, OVARC1001381, OVARC1001391, OVARC1001453, OVARC1001476, OVARC1001506, OVARC1001610, OVARC1001702, OVARC1001703, OVARC1001713, OVARC1001745, OVARC1001767, OVARC1002127, OVARC1002158, OVARC1002165, PLACE1000014, PLACE1000401, 【0991】PLACE1000562, PLACE1000611, PLACE1000656, PLACE1000712, PLACE1000909, PLACE1000948, PLACE1001241, PLACE1001257, PLACE1001377, PLACE100151

2761

7, PLACE1001610, PLACE1001771, PLACE1001817, PLACE1001983, PLACE1002213, PLACE1002395, PLACE1002500, PLACE1002714, PLACE1002722, PLACE1002794, PLACE1002851, PLACE1002908, PLACE1003045, PLACE1003238, PLACE1003296, PLACE1003369, PLACE1003493, PLACE1003537, PLACE1003553, PLACE1003768, PLACE1003771, PLACE1003903, PLACE1004197, PLACE1004258, PLACE1004270, PLACE1004289, PLACE1004473, PLACE1004743, PLACE1004840, PLACE1004969, PLACE1005086, PLACE1005206, PLACE1005313, PLACE1005530, PLACE1005595, PLACE1005623, PLACE1005763, PLACE1005884, PLACE1005934, PLACE1006225, PLACE1006754, PLACE1006901, PLACE1006935, PLACE1006956, PLACE1007014, PLACE1007111, PLACE1007243, PLACE1007274, PLACE1007282, PLACE1007317, PLACE1007375, PLACE1007386, PLACE1007409, PLACE1007484, PLACE1007583, PLACE1007632, PLACE1007645, PLACE1007852, PLACE1007877, PLACE1008331, PLACE1008424, PLACE1008531, PLACE1008532, PLACE1008568, PLACE1008715, PLACE1009045, PLACE1009319, PLACE1009338, PLACE1009368, PLACE1009493, PLACE1009639, PLACE1009708, PLACE1009731, PLACE1010089, PLACE1010231, PLACE1010321, PLACE1010622, PLACE1010811, PLACE1010917, PLACE1010954, PLACE1011090, PLACE1011214, PLACE1011221, PLACE1011399, PLACE1011492, PLACE1011646, PLACE1011749, PLACE2000034, PLACE2000111, PLACE2000176,

【0992】 PLACE2000187, PLACE2000341, PLACE2000379, PLACE2000425, PLACE2000458, PLACE3000020, PLACE3000218, PLACE3000226, PLACE3000244, PLACE3000413, PLACE4000052, PLACE4000129, PLACE4000300, PLACE4000387, PLACE4000581, PLACE4000593, PLACE4000650, THYR01000394, THYR01000395, THYR01000570, THYR01000748, THYR01000756, THYR01001134, THYR01001271, THYR01001401, THYR01001534, THYR01001541, THYR01001809, Y79AA1000258, Y79AA1000420, Y79AA1000469, Y79AA1000734, Y79AA1000800, Y79AA1000976, Y79AA1001023, Y79AA1001177, Y79AA1001394, Y79AA1001603, Y79AA1001647, Y79AA1001846, Y79AA1001874, Y79AA1002139, Y79AA1002351, Y79AA1002399, Y79AA1002416

HEMBA1004055, HEMBB1001630, HEMBB1001872, HEMBB1002044, HEMBB1002383, MAMMA1000778, MAMMA1000859, MAMMA1000897, MAMMA1001073, MAMMA1002009, MAMMA1002844, MAMMA1002947, MAMMA1003089, NT2RM1000092, NT2RM1000833, NT2RP2002105, NT2RP2003668, NT2RP2006184, NT2RP3001282, NT2RP3002810, NT2RP3002985, NT2RP3003059, NT2RP3003576, NT2RP3003665, NT2RP3003799, NT2RP3003828, NT2RP3003992, NT2RP3004051, NT2RP3004155, OVARC1000890, OVARC1001117, OVARC1001329, PLACE1001761, PLACE1002437, PLACE1004793, PLACE1005611, PLACE1005898, PLACE1009935, PLACE1011896, PLACE

2762

E2000132, PLACE2000335, PLACE3000373, PLACE3000406, PLACE4000250, PLACE4000487, PLACE4000494, THYR01001320, THYR01001537, THYR01001828, Y79AA1001384

【0993】 Pfamにより推定アミノ酸配列に機能ドメインを検出されたクローン名とヒットした機能ドメイン名は以下の通りであった。検索結果は、クローン名//機能ドメイン名のように示した。複数の機能ドメインがヒットした場合には//で区切って並記した。

HEMBA1000005//DnaJ, prokaryotic heat shock protein

HEMBA1000020//Tubulin

HEMBA1000129//Helicases conserved C-terminal domain

HEMBA1000156//RNA recognition motif. (aka RRM, RBD, or RNP domain)

HEMBA1000158//Fork head domain, eukaryotic transcription factors //Zincfinger, C2H2 type

HEMBA1000303//Src homology domain 3 //Zinc finger, C3HC4 type (RING finger)

HEMBA1000411//Ank repeat

HEMBA1000491//Ras family (contains ATP/GTP binding P-loop)

HEMBA1000531//Heat shock hsp70 proteins

HEMBA1000561//Zinc finger, C2H2 type

HEMBA1000608//Src homology domain 3

HEMBA1000919//WD domain, G-beta repeats

HEMBA1001043//Ank repeat

HEMBA1001088//LIM domain containing proteins

HEMBA1001137//Zinc finger, C2H2 type

HEMBA1001174//ADP-ribosylation factors (Arf family) (contains ATP/GTP binding P-loop)

HEMBA1001247//WW/rsp5/WWP domain containing proteins

HEMBA1001286//Sushi domain

HEMBA1001510//Basic region plus leucine zipper transcription factors

HEMBA1001515//Reverse transcriptase (RNA-dependent DNA polymerase)

HEMBA1001661//Cadherin

HEMBA1001723//WD domain, G-beta repeats

HEMBA1001744//Eukaryotic protein kinase domain

HEMBA1001804//Zinc finger, C2H2 type

HEMBA1001819//Zinc finger, C2H2 type

HEMBA1001847//Zinc finger, C2H2 type

HEMBA1002035//Bromodomain

HEMBA1002102//Ank repeat

HEMBA1002161//Myosin head (motor domain) (contains ATP/GTP binding P-loop)

HEMBA1002177//GATA family of transcription factors //Zinc finger, C2H2 type

HEMBA1002212//Eukaryotic protein kinase domain

2763

2764

HEMBA1002215//LIM domain containing proteins
 HEMBA1002419//RNA recognition motif. (aka RRM, RB
 D, or RNP domain)
 HEMBA1002547//Kazal-type serine protease inhibitor
 domain //Laminin EGF-like (Domains III and V)
 HEMBA1002768//Src homology domain 3
 HEMBA1002810//WW/rsp5/WWP domain containing protei
 ns
 HEMBA1002818//EGF-like domain
 HEMBA1002935//Zinc finger, C2H2 type
 HEMBA1002939//Ank repeat
 HEMBA1002973//3'5'-cyclic nucleotide phosphodiester
 ases
 HEMBA1003077//Fibronectin type III domain
 HEMBA1003250//Eukaryotic protein kinase domain
 HEMBA1003257//Zinc finger, C2H2 type
 HEMBA1003281//IG superfamily
 HEMBA1003291//Eukaryotic protein kinase domain
 HEMBA1003433//Forkhead-associated (FHA) domain
 HEMBA1003545//Homeobox domain //LIM domain contain
 ing proteins
 HEMBA1003591//RNA recognition motif. (aka RRM, RB
 D, or RNP domain)
 HEMBA1003684//Zinc finger, C2H2 type
 HEMBA1003953//Zinc finger, C2H2 type
 HEMBA1004202//Ras family (contains ATP/GTP binding
 P-loop)
 HEMBA1004227//Protein phosphatase 2C
 HEMBA1004321//Zinc finger, C2H2 type
 HEMBA1004356//RNA recognition motif. (aka RRM, RB
 D, or RNP domain)
 HEMBA1004408//Peptidyl-prolyl cis-trans isomerases
 HEMBA1004596//RNA recognition motif. (aka RRM, RB
 D, or RNP domain)
 HEMBA1004734//Ubiquitin-conjugating enzymes
 HEMBA1004973//Fibronectin type III domain
 HEMBA1005009//Actins
 HEMBA1005101//RNA recognition motif. (aka RRM, RB
 D, or RNP domain)
 HEMBA1005581//EGF-like domain //Laminin G domain
 HEMBA1005732//Polyprenyl synthetases
 HEMBA1005737//EF hand
 HEMBA1006248//Zinc finger, C2H2 type
 HEMBA1006284//Ubiquitin family
 HEMBA1006293//IG superfamily
 HEMBA1006344//Band 4.1 family
 HEMBA1006445//Ras family (contains ATP/GTP binding
 P-loop)
 HEMBA1006492//Ank repeat
 HEMBA1006559//Zinc finger, C3HC4 type (RING finger)

r)
 HEMBA1006708//WD domain, G-beta repeats
 HEMBA1006737//Ank repeat
 HEMBA1006758//Cadherin
 HEMBA1006941//Thioredoxins
 HEMBA1007243//Purine/pyrimidine phosphoribosyl tra
 nsferases
 HEMBA1007300//3'5'-cyclic nucleotide phosphodiester
 ases
 HEMBB1000083//IG superfamily
 HEMBB1000317//EGF-like domain //Thrombospondin typ
 e 1 domain
 HEMBB1000556//Actinin-type actin-binding domain co
 ntaining proteins //LIM domain containing proteins
 HEMBB1000725//Ras family (contains ATP/GTP binding
 P-loop)
 HEMBB1000781//Eukaryotic protein kinase domain
 HEMBB1000915//Thrombospondin type 1 domain
 HEMBB1000927//EF hand
 HEMBB1000947//Double-stranded RNA binding motif
 HEMBB1001112//eubacterial secY protein
 HEMBB1001175//Ank repeat
 HEMBB1001234//WW/rsp5/WWP domain containing protei
 ns
 HEMBB1001282//Ank repeat
 HEMBB1001294//Ras family (contains ATP/GTP binding
 P-loop)
 HEMBB1001339//Forkhead-associated (FHA) domain
 HEMBB1001673//Forkhead-associated (FHA) domain //Z
 inc finger, C3HC4 type(RING finger)
 HEMBB1001802//Intermediate filament proteins
 HEMBB1001839//Zinc finger, C2H2 type
 HEMBB1002217//Zinc finger, C2H2 type
 HEMBB1002342//Thioredoxins
 HEMBB1002600//4 transmembrane segments integral me
 mbrane proteins
 MAMMA1000173//Src homology domain 3
 MAMMA1000388//Zinc finger, C2H2 type
 MAMMA1000402//Reverse transcriptase (RNA-dependent
 DNA polymerase)
 MAMMA1000612//WD domain, G-beta repeats
 MAMMA1000672//Serine carboxypeptidases
 MAMMA1000731//SNF2 and others N-terminal domain
 MAMMA1001008//Eukaryotic aspartyl proteases
 MAMMA1001041//Actinin-type actin-binding domain co
 ntaining proteins
 MAMMA1001059//DEAD and DEAH box helicases //Helica
 ses conserved C-terminal domain
 MAMMA1001105//Zinc finger, C2H2 type
 MAMMA1001260//Zinc finger, C3HC4 type (RING finger)

- r)
- MAMMA1001576//Tubulin
- MAMMA1001735//Tubulin
- MAMMA1001768//ATPases associated with various cellular activities (AAA)
- MAMMA1001837//Zinc finger, C2H2 type
- MAMMA1002170//Ribosomal protein S5
- MAMMA1002385//RNA recognition motif. (aka RRM, RBD, or RNP domain)
- MAMMA1002619//Ubiquitin carboxyl-terminal hydrolases family 2
- MAMMA1002637//Kinesin light chain repeat
- MAMMA1002650//Zinc finger, C2H2 type
- MAMMA1002671//AMP-binding enzymes
- MAMMA1002869//LIM domain containing proteins
- MAMMA1002881//SCP-like extracellular Proteins
- MAMMA1002937//Zinc finger, C2H2 type
- MAMMA1002938//Multicopper oxidases
- MAMMA1003011//Core histones H2A, H2B, H3 and H4
- MAMMA1003057//WD domain, G-beta repeats
- MAMMA1003127//Myosin head (motor domain) (contains ATP/GTP binding P-loop)
- NT2RM1000086//Zinc finger, C3HC4 type (RING finger)
- NT2RM1000199//CUB domain //Sushi domain
- NT2RM1000256//Glutamine amidotransferases class-II
- NT2RM1000499//Ank repeat
- NT2RM1000555//'Cold-shock' DNA-binding domain containing proteins
- NT2RM1000666//'Cold-shock' DNA-binding domain containing proteins //Zincfinger, CCHC class
- NT2RM1000772//WD domain, G-beta repeats
- NT2RM1000826//'Cold-shock' DNA-binding domain containing proteins
- NT2RM1000850//Ank repeat //Eukaryotic protein kinase domain
- NT2RM1000852//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
- NT2RM1000882//Heme-binding domain in cytochrome b5 and oxidoreductases
- NT2RM1000885//Zinc finger, C3HC4 type (RING finger)
- NT2RM1001059//RNA recognition motif. (aka RRM, RBD, or RNP domain)
- NT2RM1001072//C2 domain //Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-specific phospholipase C, Y domain
- NT2RM2000092//Ubiquitin carboxyl-terminal hydrolases family 2
- NT2RM2000101//Zinc finger, C3HC4 type (RING finger)
- r)
- NT2RM2000191//3' 5'-cyclic nucleotide phosphodiesterases
- NT2RM2000422//Sodium: neurotransmitter symporter family
- NT2RM2000490//C2 domain
- NT2RM2000566//Integrins alpha chain
- NT2RM2000577//tRNA synthetases class I
- NT2RM2000594//C-5 cytosine-specific DNA methylases
- NT2RM2000691//Actins
- NT2RM2000735//Zinc finger, C2H2 type
- NT2RM2000740//Helicases conserved C-terminal domain
- NT2RM2000951//FGFY family of carbohydrate kinases
- NT2RM2001324//LIM domain containing proteins
- NT2RM2001499//Amino acid permeases
- NT2RM2001547//DnaJ, prokaryotic heat shock protein //Thioredoxins
- NT2RM2001613//eubacterial secY protein
- NT2RM2001670//Zinc finger, C2H2 type
- NT2RM2001700//Acyl-CoA dehydrogenases
- NT2RM2001730//Ubiquitin carboxyl-terminal hydrolases family 2
- NT2RM2001813//WD domain, G-beta repeats
- NT2RM2001823//Helicases conserved C-terminal domain //SNF2 and others N-terminal domain
- NT2RM2001896//Cytochrome C oxidase subunit II
- NT2RM2001989//RNA recognition motif. (aka RRM, RBD, or RNP domain)
- NT2RM2001997//Thioredoxins
- NT2RM2002088//KH domain family of RNA binding proteins
- NT2RM2002100//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
- NT2RM2002109//IG superfamily
- NT2RM4000046//Zinc finger, C3HC4 type (RING finger)
- NT2RM4000104//Zinc finger, C2H2 type
- NT2RM4000167//Kinesin motor domain
- NT2RM4000191//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
- NT2RM4000202//Zinc finger, C2H2 type
- NT2RM4000229//PH (pleckstrin homology) domain
- NT2RM4000344//ATPases associated with various cellular activities (AAA)
- NT2RM4000356//Ras family (contains ATP/GTP binding P-loop)
- NT2RM4000471//Aminotransferases class-V
- NT2RM4000496//ATPases associated with various cellular activities (AAA)

2767

NT2RM4000611//WD domain, G-beta repeats
 NT2RM4000657//C2 domain //Phosphatidylinositol-specific phospholipase C,Y domain
 NT2RM4000712//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RM4000733//Forkhead-associated (FHA) domain
 NT2RM4000734//Zinc finger, C2H2 type
 NT2RM4000751//Zinc finger, C2H2 type
 NT2RM4000795//Carboxylesterases
 NT2RM4000996//Zinc finger, C2H2 type
 NT2RM4001054//eubacterial secY protein
 NT2RM4001140//Homeobox domain
 NT2RM4001178//DEAD and DEAH box helicases
 NT2RM4001200//Zinc finger, C2H2 type
 NT2RM4001313//Phosphatidylinositol 3- and 4-kinases
 NT2RM4001316//Acyl-CoA dehydrogenases
 NT2RM4001320//Src homology domain 3
 NT2RM4001411//PH (pleckstrin homology) domain //Src homology domain 2
 NT2RM4001454//PH (pleckstrin homology) domain
 NT2RM4001483//Zinc finger, C2H2 type
 NT2RM4001629//Src homology domain 3
 NT2RM4001758//Eukaryotic protein kinase domain
 NT2RM4001810//Zinc finger, C2H2 type
 NT2RM4001813//Lectin C-type domain short and long forms
 NT2RM4001823//Zinc finger, C2H2 type
 NT2RM4001828//Zinc finger, C2H2 type
 NT2RM4001979//Zinc finger, C2H2 type
 NT2RM4001987//IG superfamily
 NT2RM4002013//WD domain, G-beta repeats
 NT2RM4002073//AMP-binding enzymes
 NT2RM4002093//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RM4002145//IG superfamily
 NT2RM4002287//Fibronectin type III domain
 NT2RM4002527//WD domain, G-beta repeats
 NT2RM4002623//tRNA synthetases class II
 NT2RP1000101//Zinc finger, C2H2 type
 NT2RP1000202//Ank repeat
 NT2RP1000272//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP1000363//PH (pleckstrin homology) domain
 NT2RP1000376//Ank repeat
 NT2RP1000470//DEAD and DEAH box helicases
 NT2RP1000478//Tubulin
 NT2RP1000522//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases

2768

es family 2
 NT2RP1000677//Kazal-type serine protease inhibitor domain
 NT2RP1000701//WD domain, G-beta repeats
 NT2RP1000733//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP1000782//4 transmembrane segments integral membrane proteins
 NT2RP1000833//3' 5'-cyclic nucleotide phosphodiesterases
 NT2RP1000856//4 transmembrane segments integral membrane proteins
 NT2RP1000947//Ubiquitin-conjugating enzymes
 NT2RP1000959//60s Acidic ribosomal protein
 NT2RP1000966//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP1001033//Tubulin
 NT2RP1001080//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP1001177//Core histones H2A, H2B, H3 and H4
 NT2RP1001247//Transforming growth factor beta like domain
 NT2RP1001294//WD domain, G-beta repeats
 NT2RP1001302//WD domain, G-beta repeats
 NT2RP1001313//Heme-binding domain in cytochrome b5 and oxidoreductases
 NT2RP1001457//WD domain, G-beta repeats
 NT2RP1001546//4 transmembrane segments integral membrane proteins
 NT2RP2000008//Zinc finger, C2H2 type
 NT2RP2000040//C2 domain
 NT2RP2000045//DnaJ, prokaryotic heat shock protein
 NT2RP2000054//Zinc finger, C3HC4 type (RING finger)
 NT2RP2000070//Cadherin
 NT2RP2000126//Helicases conserved C-terminal domain //SNF2 and others N-terminal domain
 NT2RP2000153//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP2000224//PH (pleckstrin homology) domain
 NT2RP2000257//Mitochondrial carrier proteins
 NT2RP2000329//Adenylate kinases
 NT2RP2000414//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP2000448//PH (pleckstrin homology) domain
 NT2RP2000660//ATPases associated with various cellular activities (AAA)
 NT2RP2000668//Eukaryotic protein kinase domain
 NT2RP2000710//tRNA synthetases class II
 NT2RP2000764//Aminotransferases class-V

- NT2RP2000842//7 transmembrane receptor (rhodopsin family)
 NT2RP2000880//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP2000931//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP2000932//Ank repeat
 NT2RP2001081//C2 domain
 NT2RP2001174//Zinc finger, C2H2 type
 NT2RP2001397//Cyclins
 NT2RP2001520//Mitochondrial carrier proteins
 NT2RP2001597//Zinc finger, C3HC4 type (RING finger)
 NT2RP2001740//Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP2001748//Polyprenyl synthetases
 NT2RP2001756//Zinc finger, C2H2 type
 NT2RP2001839//Eukaryotic protein kinase domain
 NT2RP2001900//Actins
 NT2RP2001991//Sodium:neurotransmitter symporter family
 NT2RP2002058//WD domain, G-beta repeats
 NT2RP2002124//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP2002185//Ubiquitin family
 NT2RP2002208//Zinc finger, C3HC4 type (RING finger)
 NT2RP2002256//Cytochrome P450
 NT2RP2002479//ABC transporters
 NT2RP2002503//Zinc finger, C2H2 type
 NT2RP2002520//Ank repeat
 NT2RP2002591//Zinc finger, C2H2 type
 NT2RP2002741//Src homology domain 3
 NT2RP2002929//WD domain, G-beta repeats
 NT2RP2002939//Zinc finger, C2H2 type
 NT2RP2002959//Ubiquitin-conjugating enzymes
 NT2RP2002980//Ribosomal protein S10
 NT2RP2003137//Ubiquitin family
 NT2RP2003164//Eukaryotic protein kinase domain
 NT2RP2003228//MC2/3/5 family
 NT2RP2003243//Fibronectin type III domain
 NT2RP2003272//Ubiquitin family
 NT2RP2003307//Kinesin light chain repeat
 NT2RP2003401//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP2003433//eubacterial secY protein
 NT2RP2003480//Zinc finger, C2H2 type
 NT2RP2003713//Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP2003737//Ubiquitin-conjugating enzymes
 NT2RP2003777//Zinc finger, C3HC4 type (RING finger)
 NT2RP2003840//Ubiquitin-conjugating enzymes
 NT2RP2003857//Ank repeat
 NT2RP2003981//Zinc finger, C3HC4 type (RING finger)
 NT2RP2004170//WD domain, G-beta repeats
 10 NT2RP2004187//Zinc finger, C2H2 type
 NT2RP2004232//Phorbol esters / diacylglycerol binding domain //PH (pleckstrin homology) domain //Eukaryotic protein kinase domain
 NT2RP2004389//Ribosomal protein S9
 NT2RP2004538//PH (pleckstrin homology) domain
 NT2RP2004568//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP2004710//WW/rsp5/WWP domain containing proteins
 NT2RP2004768//Eukaryotic protein kinase domain
 NT2RP2004933//Eukaryotic protein kinase domain
 NT2RP2004961//Zinc finger, C2H2 type
 NT2RP2005003//Zinc finger, C3HC4 type (RING finger)
 NT2RP2005012//DnaJ, prokaryotic heat shock protein
 NT2RP2005126//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP2005139//Ank repeat
 NT2RP2005140//PH (pleckstrin homology) domain
 30 NT2RP2005239//Aminotransferases class-V
 NT2RP2005288//Regulator of chromosome condensation (RCC1)
 NT2RP2005293//PH (pleckstrin homology) domain
 NT2RP2005325//Homeobox domain //LIM domain containing proteins
 NT2RP2005344//E1-E2 ATPases
 NT2RP2005465//Mitochondrial carrier proteins
 NT2RP2005525//Forkhead-associated (FHA) domain
 NT2RP2005531//Band 4.1 family
 40 NT2RP2005557//Bacterial mutT protein
 NT2RP2005654//DnaJ, prokaryotic heat shock protein
 NT2RP2005701//Zinc finger, C3HC4 type (RING finger)
 NT2RP2005722//Zinc finger, C2H2 type
 NT2RP2005752//TNFR/NGFR cysteine-rich region
 NT2RP2005763//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP2005767//HMG (high mobility group) box
 NT2RP2006312//HMG (high mobility group) box
 50 NT2RP2006464//HMG (high mobility group) box

2771

NT2RP2006571//Cytochrome P450
 NT2RP3000050//Zinc finger, C2H2 type
 NT2RP3000068//PH (pleckstrin homology) domain
 NT2RP3000085//Biotin-requiring enzymes //Carbamoyl-phosphate synthase (CPSase)
 NT2RP3000299//Src homology domain 3
 NT2RP3000359//Adenylate kinases
 NT2RP3000366//Ras family (contains ATP/GTP binding P-loop)
 NT2RP3000403//WW/rsp5/WWP domain containing proteins
 NT2RP3000487//WW/rsp5/WWP domain containing proteins
 NT2RP3000512//Homeobox domain
 NT2RP3000527//Zinc finger, C2H2 type
 NT2RP3000531//IG superfamily
 NT2RP3000590//Zinc finger, C3HC4 type (RING finger)
 NT2RP3000603//Helix-loop-helix DNA-binding domain
 NT2RP3000605//Zinc finger, C2H2 type
 NT2RP3000632//Zinc finger, C2H2 type
 NT2RP3000742//Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-specific phospholipase C, Y domain
 NT2RP3000759//ADP-ribosylation factors (Arf family) (contains ATP/GTP binding P-loop)
 NT2RP3000825//EGF-like domain
 NT2RP3000869//ATPases associated with various cellular activities (AAA)
 NT2RP3000994//Double-stranded RNA binding motif
 NT2RP3001057//Zinc finger, C2H2 type
 NT2RP3001084//PH (pleckstrin homology) domain
 NT2RP3001120//Zinc finger, C2H2 type
 NT2RP3001140//Thrombospondin type 1 domain
 NT2RP3001150//Forkhead-associated (FHA) domain
 NT2RP3001155//HMG (high mobility group) box
 NT2RP3001214//Zinc finger, C2H2 type
 NT2RP3001268//Zinc finger, C2H2 type
 NT2RP3001338//Zinc finger, C2H2 type
 NT2RP3001355//Mitochondrial carrier proteins
 NT2RP3001398//Zinc finger, C2H2 type
 NT2RP3001426//DnaJ, prokaryotic heat shock protein
 NT2RP3001453//ABC transporters
 NT2RP3001457//PH (pleckstrin homology) domain
 NT2RP3001472//HMG (high mobility group) box
 NT2RP3001495//Alcohol/other dehydrogenases, short chain type //WW/rsp5/WWP domain containing proteins
 NT2RP3001497//Zinc finger, C3HC4 type (RING finger)

2772

NT2RP3001724//Helicases conserved C-terminal domain
 NT2RP3001792//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP3001943//Zinc finger, C3HC4 type (RING finger)
 NT2RP3001944//Zinc finger, C3HC4 type (RING finger)
 NT2RP3002007//ATPases associated with various cellular activities (AAA)
 NT2RP3002054//Low-density lipoprotein receptor domain class A
 NT2RP3002151//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP3002399//MCM2/3/5 family
 NT2RP3002501//Serine/threonine dehydratases
 NT2RP3002602//Thioredoxins
 NT2RP3002628//DnaJ, prokaryotic heat shock protein //Thioredoxins
 20 NT2RP3002663//PH (pleckstrin homology) domain
 NT2RP3002909//Ank repeat
 NT2RP3002953//Cadherin
 NT2RP3002969//AMP-binding enzymes
 NT2RP3003061//Ank repeat
 NT2RP3003145//Zinc carboxypeptidases
 NT2RP3003230//WD domain, G-beta repeats
 NT2RP3003251//Zinc finger, C3HC4 type (RING finger)
 NT2RP3003278//Ank repeat //Zinc finger, C2H2 type
 30 NT2RP3003282//PH (pleckstrin homology) domain
 NT2RP3003311//PH (pleckstrin homology) domain
 NT2RP3003385//Ank repeat //Chaperonins clpA/B
 NT2RP3003589//Ras family (contains ATP/GTP binding P-loop)
 NT2RP3003621//CUB domain //Kringle domain
 NT2RP3003701//Thrombospondin type 1 domain
 NT2RP3003716//Fibronectin type III domain
 NT2RP3003809//ATPases associated with various cellular activities (AAA)
 40 NT2RP3004016//Zinc finger, C3HC4 type (RING finger)
 NT2RP3004207//CUB domain //Sushi domain
 NT2RP3004209//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP3004242//PH (pleckstrin homology) domain
 NT2RP3004262//DnaJ, prokaryotic heat shock protein
 NT2RP3004566//Zinc finger, C2H2 type
 NT2RP3004569//Ank repeat
 50 NT2RP3004594//HMG (high mobility group) box

2773

NT2RP3004617//Zinc finger, C3HC4 type (RING finger)
 NT2RP4000259//Glutathione peroxidases
 NT2RP4000370//Prokaryotic-type class I peptide chain release factors
 NT2RP4000376//WD domain, G-beta repeats
 NT2RP4000398//Zinc finger, C2H2 type
 NT2RP4000455//Forkhead-associated (FHA) domain //Zinc finger, C3HC4 type (RING finger)
 NT2RP4000457//Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP4000518//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP4000588//Actinin-type actin-binding domain containing proteins
 NT2RP4000614//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP4000648//Forkhead-associated (FHA) domain //Zinc finger, C3HC4 type (RING finger)
 NT2RP4000837//Zinc finger, C2H2 type
 NT2RP4000839//WD domain, G-beta repeats
 NT2RP4000865//Zinc finger, C2H2 type
 NT2RP4000907//Fibronectin type III domain //IG superfamily
 NT2RP4000925//Fibronectin type III domain
 NT2RP4000927//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP4000973//DnaJ, prokaryotic heat shock protein //Thioredoxins
 NT2RP4001079//E1-E2 ATPases
 NT2RP4001080//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP4001117//eubacterial secY protein
 NT2RP4001150//Fibronectin type III domain
 NT2RP4001213//Zinc finger, C2H2 type
 NT2RP4001219//Thioredoxins
 NT2RP4001235//Zinc finger, CCHC class
 NT2RP4001433//Zinc finger, C2H2 type
 NT2RP4001498//Ank repeat
 NT2RP4001568//Ank repeat
 NT2RP4001644//Eukaryotic protein kinase domain
 NT2RP4001725//WD domain, G-beta repeats
 NT2RP4001753//Zinc finger, C2H2 type
 NT2RP4001790//Zinc finger, C2H2 type
 NT2RP4001822//4 transmembrane segments integral membrane proteins
 NT2RP4001823//Fibrinogen beta and gamma chains, C-terminal globular domain
 NT2RP4001893//Ank repeat

2774

NT2RP4001896//WD domain, G-beta repeats
 NT2RP4001927//WD domain, G-beta repeats
 NT2RP4001938//Zinc finger, C2H2 type
 NT2RP4002047//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP4002078//Zinc finger, C2H2 type
 NT2RP4002408//Eukaryotic protein kinase domain
 NT2RP4002905//Cyclins
 NT2RP5003477//WD domain, G-beta repeats
 OVARC1000006//Core histones H2A, H2B, H3 and H4
 OVARC1000085//Proteasome A-type and B-type
 OVARC1000148//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 OVARC1000556//Eukaryotic protein kinase domain
 OVARC1000649//PH (pleckstrin homology) domain //Src homology domain 2
 OVARC1000746//Double-stranded RNA binding motif
 OVARC1000885//Alcohol/other dehydrogenases, short chain type
 OVARC1000937//Cyclins
 OVARC1000999//Ank repeat
 OVARC1001154//Granulins
 OVARC1001180//Ubiquitin family
 OVARC1001306//Helix-loop-helix DNA-binding domain
 OVARC1001577//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 OVARC1001731//Tropomyosins
 OVARC1001943//Zinc finger, C2H2 type
 OVARC1002050//Spectrin alpha chain, repeated domain
 OVARC1002112//Core histones H2A, H2B, H3 and H4
 OVARC1002138//ATPases associated with various cellular activities (AAA)
 OVARC1002182//WD domain, G-beta repeats
 PLACE1000014//Zinc finger, C3HC4 type (RING finger)
 PLACE1000040//Ras family (contains ATP/GTP binding P-loop)
 PLACE1000050//Zinc finger, C2H2 type
 PLACE1000081//PH (pleckstrin homology) domain
 PLACE1000142//Enoyl-CoA hydratase/isomerase
 PLACE1000401//IG superfamily
 PLACE1000406//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1000420//Bacterial mutT protein
 PLACE1000706//Bromodomain
 PLACE1000769//KH domain family of RNA binding proteins
 PLACE1000786//PH (pleckstrin homology) domain
 PLACE1000863//Ribosomal protein S4

2775

PLACE1000909//Ank repeat
 PLACE1000972//Src homology domain 3
 PLACE1000979//Zinc finger, C2H2 type
 PLACE1001304//Zinc finger, C2H2 type
 PLACE1001387//Src homology domain 3
 PLACE1001632//Zinc finger, C2H2 type
 PLACE1001672//Aminotransferases class-III pyridoxal-phosphate
 PLACE1001716//Zinc finger, CCHC class
 PLACE1001739//DEAD and DEAH box helicases //Helices conserved C-terminal domain
 PLACE1001781//Phosphoglucosyltransferase and phosphomannosyltransferase phosphoserine
 PLACE1001869//FGFY family of carbohydrate kinases
 PLACE1002438//Zinc finger, C2H2 type
 PLACE1002450//Zinc finger, C2H2 type
 PLACE1002474//EGF-like domain //von Willebrand factor type A domain
 PLACE1002499//Zinc finger, C3HC4 type (RING finger)
 PLACE1002532//Homeobox domain
 PLACE1002571//Actins
 PLACE1002685//Src homology domain 2
 PLACE1002722//7 transmembrane receptor (rhodopsin family)
 PLACE1002775//Bromodomain
 PLACE1002834//Zinc finger, C2H2 type
 PLACE1003100//Alcohol/other dehydrogenases, short chain type
 PLACE1003174//Ubiquitin-conjugating enzymes
 PLACE1003238//7 transmembrane receptor (rhodopsin family)
 PLACE1003302//Zinc finger, C2H2 type
 PLACE1003334//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1003366//C2 domain
 PLACE1003394//Ras family (contains ATP/GTP binding P-loop)
 PLACE1003420//Mitochondrial carrier proteins
 PLACE1003493//Clq domain
 PLACE1003519//KH domain family of RNA binding proteins
 PLACE1003723//Src homology domain 2
 PLACE1003738//Zinc finger, C2H2 type
 PLACE1003888//C2 domain //Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-specific phospholipase C, Y domain
 PLACE1004128//WD domain, G-beta repeats
 PLACE1004358//PH (pleckstrin homology) domain
 PLACE1004428//Acyl-CoA dehydrogenases

2776

PLACE1004437//Isocitrate and isopropylmalate dehydrogenases
 PLACE1004506//LIM domain containing proteins
 PLACE1004674//EF hand
 PLACE1004918//L-lactate dehydrogenases
 PLACE1005243//Eukaryotic protein kinase domain
 PLACE1005305//Adenylate kinases
 PLACE1005327//Src homology domain 3
 PLACE1005530//Zinc finger, C3HC4 type (RING finger)
 PLACE1005646//Helicases conserved C-terminal domain
 PLACE1005656//Ribonucleotide reductases
 PLACE1005966//WD domain, G-beta repeats
 PLACE1006157//Sushi domain
 PLACE1006196//DEAD and DEAH box helicases //Helices conserved C-terminal domain
 PLACE1006438//Zinc finger, C2H2 type
 PLACE1006626//Double-stranded RNA binding motif
 PLACE1006754//IG superfamily
 PLACE1006829//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 PLACE1006917//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1006956//ABC transporters
 PLACE1006958//Heat shock hsp70 proteins
 PLACE1007375//C2 domain
 PLACE1007488//PH (pleckstrin homology) domain
 PLACE1007511//Intermediate filament proteins
 PLACE1007537//Ank repeat
 PLACE1007544//Zinc finger, C2H2 type
 PLACE1007547//Zinc finger, C3HC4 type (RING finger)
 PLACE1007598//Zinc finger, C2H2 type
 PLACE1007697//ABC transporters
 PLACE1007958//3' 5'-cyclic nucleotide phosphodiesterases
 PLACE1007969//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1008201//Zinc finger, C2H2 type
 PLACE1008429//Ank repeat
 PLACE1008465//Zinc finger, C2H2 type
 PLACE1008650//WD domain, G-beta repeats
 PLACE1009020//Aminotransferases class-V
 PLACE1009094//von Willebrand factor type C domain
 PLACE1009099//Zinc finger, C2H2 type
 PLACE1009246//LIM domain containing proteins
 PLACE1009468//WD domain, G-beta repeats
 PLACE1009476//DEAD and DEAH box helicases //Helices conserved C-terminal domain

2777

ses conserved C-terminal domain
 PLACE1009524//PH (pleckstrin homology) domain
 PLACE1009596//WD domain, G-beta repeats
 PLACE1009622//Double-stranded RNA binding motif
 PLACE1009861//Cysteine proteases
 PLACE1009925//Helicases conserved C-terminal domain
 PLACE1009992//CUB domain //EGF-like domain //Sushi domain //Trypsin
 PLACE1010053//Double-stranded RNA binding motif
 PLACE1010089//Ubiquitin carboxyl-terminal hydrolases family 2
 PLACE1010702//Zinc finger, C2H2 type
 PLACE1010833//EF hand
 PLACE1010926//Src homology domain 3
 PLACE1010960//Actins
 PLACE1011041//Src homology domain 3
 PLACE1011046//C2 domain //Phosphatidylinositol-specific phospholipase C,X domain //Phosphatidylinositol-specific phospholipase C, Y domain
 PLACE1011114//Helicases conserved C-terminal domain
 PLACE1011160//Zinc finger, C3HC4 type (RING finger)
 PLACE1011263//Ank repeat
 PLACE1011433//Zinc finger, C2H2 type
 PLACE1011576//Zinc finger, C2H2 type
 PLACE1011923//Eukaryotic protein kinase domain
 PLACE2000034//Fibronectin type III domain //IG superfamily
 PLACE2000072//Zinc finger, C2H2 type
 PLACE2000111//IG superfamily
 PLACE2000164//WD domain, G-beta repeats
 PLACE2000216//PH (pleckstrin homology) domain
 PLACE2000341//Sodium: solute symporter family
 PLACE2000371//Src homology domain 2
 PLACE2000373//Thrombospondin type 1 domain
 PLACE2000398//IG superfamily
 PLACE2000427//Helicases conserved C-terminal domain
 PLACE2000458//Cadherin
 PLACE3000020//Guanylate cyclases
 PLACE3000169//Zinc finger, C2H2 type
 PLACE4000014//Helicases conserved C-terminal domain
 PLACE4000052//ABC transporters
 PLACE4000192//Zinc finger, C2H2 type
 PLACE4000211//Bromodomain
 PLACE4000431//Helicases conserved C-terminal domain

2778

PLACE4000522//Ank repeat
 PLACE4000581//EGF-like domain //Sushi domain
 PLACE4000654//Ubiquitin-conjugating enzymes
 THYR01000072//IG superfamily
 THYR01000242//Zinc finger, C2H2 type
 THYR01000288//Zinc-binding metalloprotease domain
 THYR01000488//Zinc finger, C3HC4 type (RING finger)
 THYR01000501//Zinc finger, C3HC4 type (RING finger)
 THYR01000666//Kinesin motor domain
 THYR01000748//Src homology domain 3
 THYR01000926//3' 5'-cyclic nucleotide phosphodiesterases
 THYR01001661//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 THYR01001671//Ubiquitin family
 Y79AA1000037//Zinc finger, C3HC4 type (RING finger)
 Y79AA1000214//Core histones H2A, H2B, H3 and H4
 Y79AA1000342//Zinc finger, C2H2 type
 Y79AA1000349//Double-stranded RNA binding motif
 Y79AA1000627//Zinc finger, C2H2 type
 Y79AA1000705//Helicases conserved C-terminal domain
 Y79AA1000752//KH domain family of RNA binding proteins
 Y79AA1000833//Tubulin
 Y79AA1001048//Acyl-CoA dehydrogenases
 Y79AA1001391//Homeobox domain
 Y79AA1001394//ATPases associated with various cellular activities (AAA)
 Y79AA1001493//Ubiquitin-conjugating enzymes
 Y79AA1001613//Zinc finger, C2H2 type
 Y79AA1001874//TNFR/NGFR cysteine-rich region
 Y79AA1002027//Ubiquitin-conjugating enzymes
 Y79AA1002139//DnaJ, prokaryotic heat shock protein
 Y79AA1002208//Ank repeat
 Y79AA1002246//C2 domain
 Y79AA1002307//Fibronectin type III domain
 Y79AA1002472//Zinc finger, C2H2 type
 HEMBA1003538//CUB domain
 HEMBA1003645//WD domain, G-beta repeats //Src homology domain 3
 HEMBA1005206//Glutathione S-transferases.
 HEMBA1006521//Alcohol/other dehydrogenases, short chain type
 HEMBB1001482//Zinc finger, C2H2 type
 HEMBB1001915//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases

2779

es family 2

HEMBB1002044//Cadherin

MAMMA1000183//Zinc finger, C2H2 type

MAMMA1000897//von Willebrand factor type A domain

MAMMA1001080//IG superfamily

MAMMA1002498//IG superfamily

MAMMA1002573//KH domain family of RNA binding proteins

MAMMA1002617//Zinc finger, C2H2 type

NT2RM1000833//eubacterial secY protein

NT2RM2001797//Zinc finger, C2H2 type

NT2RP1001013//Zinc finger, C2H2 type

NT2RP2001233//Zinc finger, C2H2 type

NT2RP2001440//14-3-3 proteins

NT2RP2002105//7 transmembrane receptor (rhodopsin family)

NT2RP3001723//Laminin G domain

NT2RP3001938//Eukaryotic protein kinase domain

NT2RP3002330//Elongation factor Tu family (contains ATP/GTP binding P-loop)

NT2RP3003133//Zinc finger, C2H2 type

NT2RP3003500//Eukaryotic protein kinase domain

NT2RP3003799//C2 domain

NT2RP3003800//Eukaryotic protein kinase domain

NT2RP3004013//Double-stranded RNA binding motif

NT2RP3004125//Zinc finger, C2H2 type

OVARC1001244//Bromodomain

OVARC1001496//D-isomer specific 2-hydroxyacid dehydrogenases

PLACE1000007//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2

PLACE1001118//Zinc finger, C2H2 type

PLACE1010310//Zinc finger, C2H2 type

PLACE1011896//wnt family of developmental signaling proteins

PLACE3000124//Src homology domain 2

PLACE4000100//D-isomer specific 2-hydroxyacid dehydrogenases

PLACE4000259//Helicases conserved C-terminal domain

PLACE4000261//Bromodomain

SKNMC1000013//ABC transporters

SKNMC1000091//Basic region plus leucine zipper transcription factors

THYR01000343//Src homology domain 3

THYR01000569//Zinc finger, C2H2 type

THYR01001189//Zinc finger, C2H2 type

Y79AA1002103//Zinc finger, C2H2 type

PLACE3000350//Eukaryotic protein kinase domain

2780

PLACE4000156//Zinc finger, C2H2 type

【0994】実施例18. 全長配列による機能カテゴリー分類

全長塩基配列が明らかになった4997クローンについてGenBank、Swiss-Prot、UniGeneの各データベースを対象に行った同源性検索の結果(同源性検索結果6、12、13、14参照)や、全長塩基配列から推定されたアミノ酸配列に対するドメイン検索の結果(実施例17参照)から、クローン中にコードされるタンパク質の機能予測、カテゴリー分類を行った。対象とした4997クローンのリストは以下の通りである。

HEMBA1000005, HEMBA1000012, HEMBA1000020, HEMBA1000030, HEMBA1000042, HEMBA1000046, HEMBA1000050, HEMBA1000076, HEMBA1000129, HEMBA1000141, HEMBA1000150, HEMBA1000156, HEMBA1000158, HEMBA1000168, HEMBA1000185, HEMBA1000193, HEMBA1000201, HEMBA1000213, HEMBA1000216, HEMBA1000227, HEMBA1000231, HEMBA1000243, HEMBA1000244, HEMBA1000251, HEMBA1000264, HEMBA1000280, HEMBA1000282, HEMBA1000288, HEMBA1000290, HEMBA1000302, HEMBA1000303, HEMBA1000304, HEMBA1000307, HEMBA1000327, HEMBA1000333, HEMBA1000338, HEMBA1000351, HEMBA1000356, HEMBA1000357, HEMBA1000369, HEMBA1000376, HEMBA1000387, HEMBA1000392, HEMBA1000396, HEMBA1000411, HEMBA1000428, HEMBA1000442, HEMBA1000456, HEMBA1000459, HEMBA1000460, HEMBA1000469, HEMBA1000488, HEMBA1000491, HEMBA1000497, HEMBA1000501, HEMBA1000504, HEMBA1000505, HEMBA1000508, HEMBA1000518, HEMBA1000519, HEMBA1000520, HEMBA1000523, HEMBA1000531, HEMBA1000534, HEMBA1000542, HEMBA1000545, HEMBA1000555, HEMBA1000557, HEMBA1000561, HEMBA1000568, HEMBA1000569, 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2792

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2796

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2797

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2798

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2799

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2800

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2801

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2802

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【1043】4997クローンのうち、分泌・膜タンパク質、糖タンパク質関連タンパク質、シグナル伝達関連タンパク質、転写関連タンパク質、疾患関連タンパク質、酵素・代謝関連タンパク質、ATP・GTP結合タンパク質、核タンパク質、DNA・RNA結合タンパク質、RNA合成関連タンパク質、タンパク質合成・輸送関連タンパク質、細胞骨格関連タンパク質、細胞分裂・増殖関連タンパク質、発生、発達関連タンパク質、細胞防御関連タンパク質のいずれかのカテゴリーに属するタンパク質をコードすると推定されるクローンは2189クローンあった。

【1044】分泌・膜タンパク質のカテゴリーに属するタンパク質をコードすると推定されるクローンは、全長配列に対するSwiss-Protヒットデータのキーワード中に "growth factor", "cytokine", "hormone", "signal", "transmembrane", "membrane", "extracellular matrix", "receptor", "G-protein coupled receptor", "ionic channel", "voltage-gated channel", "calcium channel", "cell adhesion", "collagen", "connective tissue" があつた、またはヒットデータ中に分泌・膜タンパク質と推定される記載があつた、あるいは全長配列に対するGenBank、UniGeneヒットデータ中に同様の記載があつた、さらには全長塩基配列から推定されたアミノ酸配列に対するドメイン検索の結果から、N末端シグナル配列や膜貫通領域が予測されたクローンである。

【1045】糖タンパク質関連タンパク質のカテゴリーに属するタンパク質をコードすると推定されるクローンは、全長配列に対するSwiss-Protヒットデータのキーワード中に "glycoprotein" があつた、またはヒットデータ中に糖タンパク質と推定される記載があつた、あるいは全長配列に対するGenBank、UniGeneヒットデータ中に同様の記載があつたクローンである。

【1046】シグナル伝達関連タンパク質のカテゴリーに属するタンパク質をコードすると推定されるクローンは、全長配列に対するSwiss-Protヒットデータのキーワード中に "serine/threonine-protein kinase", "tyrosine-protein kinase", "SH3 domain", があつた、またはヒットデータ中にシグナル伝達関連タンパク質と推定される記載 (例えば、"ADP-ribosylation factor") があつた、あるいは全長配列に対するGenBank、UniGeneヒットデータ中に同様の記載があつたクローンである。

転写関連タンパク質のカテゴリーに属するタンパク質を

コードすると推定されるクローンは、全長配列に対するSwiss-Protヒットデータのキーワード中に "transcription regulation", "zinc finger", "homeobox" があつた、またはヒットデータ中に転写関連タンパク質と推定される記載があつた、あるいは全長配列に対するGenBank、UniGeneヒットデータ中に同様の記載があつたクローンである。

【1047】疾患関連タンパク質のカテゴリーに属するタンパク質をコードすると推定されるクローンは、全長配列に対するSwiss-Protヒットデータのキーワード中に "disease mutation", "syndrome" があつた、またはヒットデータ中に疾患関連タンパク質と推定される記載があつた、あるいは全長配列に対するSwiss-Protヒットデータ、及びGenBank、UniGeneヒットデータが、ヒトの遺伝子と疾患のデータベースであるOnline Mendelian Inheritance in Man (OMIM)に登録されている遺伝子、タンパク質であつたクローンである。酵素・代謝関連タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にmetabolism, oxidoreductase, E. C.No. (Enzyme commission number)等と記載があつたクローンであつた。

【1048】ATP・GTP結合タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にATP-binding, GTP-binding等と記載があつたクローンであつた。核タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にnuclear protein等と記載があつたクローンであつた。DNA・RNA結合タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にDNA-binding, RNA-binding等と記載があつたクローンであつた。RNA合成関連タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にRNA splicing, RNA processing, RNA helicase, polyadenylation等と記載があつたクローンであつた。

【1049】タンパク質合成・輸送関連タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にtranslation regulation, protein biosynthesis, amino-acid biosynthesis, ribosomal protein, protein transport, signal recognition particle等と記載があつたクローンであつた。細胞骨格関連タンパク質のカテゴリーに属すると推定されたクローンは、ヒットデータ中にstructural protein, cytoskeleton, actin-binding, microtubules等と記載があつたクローンであつた。

【1050】細胞分裂・増殖関連タンパク質のカテゴリーに属すると推定されたクローンは、cell division, cell cycle, mitosis, chromosomal protein, cell growth, apoptosis等と記載があつたクローンであつた。発生・発達関連タンパク質のカテゴリーに属すると推定されたクローンは、developmental protein等と記載があつたクローンであつた。細胞防御関連タンパク質の

2811

カテゴリーに属すると推定されたクローンとは、ヒットデータ中にheat shock, DNA repair, DNA damage等と記載があったクローンであった。この機能カテゴリー分類では一つのクローンが上記の複数のカテゴリーに該当する場合は、そのまま複数のカテゴリーに分類した。ただし、タンパク質の機能は必ずしも分類された機能カテゴリーに限定されるわけではなく、今後その他の機能も明らかになる可能性がある。

【1051】分泌・膜タンパク質に分類されたクローンは以下の796クローンであった。

HEMBA1000356, HEMBA1000518, HEMBA1000531, HEMBA100637, HEMBA1000719, HEMBA1000817, HEMBA1000822, HEMBA1000852, HEMBA1000870, HEMBA1000991, HEMBA1001052, HEMBA1001071, HEMBA1001085, HEMBA1001286, HEMBA1001351, HEMBA1001407, HEMBA1001446, HEMBA1001515, HEMBA1001557, HEMBA1001569, HEMBA1001661, HEMBA1001734, HEMBA1001746, HEMBA1001866, HEMBA1002125, HEMBA1002150, HEMBA1002166, HEMBA1002417, HEMBA1002462, HEMBA1002475, HEMBA1002477, HEMBA1002486, HEMBA1002609, HEMBA1002659, HEMBA1002661, HEMBA1002780, HEMBA1002818, HEMBA1002876, HEMBA1002921, HEMBA1003071, HEMBA1003077, HEMBA1003079, HEMBA1003086, HEMBA1003096, HEMBA1003281, HEMBA1003286, HEMBA1003538, HEMBA1003711, HEMBA1003742, HEMBA1003803, HEMBA1004055, HEMBA1004143, HEMBA1004146, HEMBA1004207, HEMBA1004341, HEMBA1004461, HEMBA1004577, HEMBA1004637, HEMBA1004752, HEMBA1004756, HEMBA1004850, HEMBA1004889, HEMBA1004923, HEMBA1004930, HEMBA1005029, HEMBA1005035, HEMBA1005050, HEMBA1005552, HEMBA1005576, HEMBA1005581, HEMBA1005588, HEMBA1005616, HEMBA1005699, HEMBA1005991, HEMBA1006036, HEMBA1006038, HEMBA1006067, HEMBA1006173, HEMBA1006198, HEMBA1006293, HEMBA1006310, HEMBA1006492, HEMBA1006502, HEMBA1006583, HEMBA1006659, HEMBA1006758, HEMBA1006789, HEMBA1006921, HEMBA1006926, HEMBA1006976, HEMBA1007203, HEMBA1007301, HEMBB1000037, HEMBB1000050, HEMBB1000054, HEMBB1000175, HEMBB1000317, HEMBB1000556, HEMBB1000593, HEMBB1000631, 【1052】HEMBB1000763, HEMBB1000827, HEMBB1000915, HEMBB1000975, HEMBB1001112, HEMBB1001151, HEMBB1001177, HEMBB1001302, HEMBB1001348, HEMBB1001564, HEMBB1001630, HEMBB1001871, HEMBB1001872, HEMBB1001925, HEMBB1001962, HEMBB1002042, HEMBB1002044, HEMBB1002142, HEMBB1002190, HEMBB1002193, HEMBB1002247, HEMBB1002383, HEMBB1002387, HEMBB1002550, HEMBB1002600, HEMBB1002692, MAMMA1000045, MAMMA1000129, MAMMA1000133, MAMMA1000277, MAMMA1000278, MAMMA1000410, MAMMA1000416, MAMMA1000472, MAMMA1000672, MAMMA1000684, MAMMA1000714, MAMMA1000734, MAMMA1000778, MAMMA1000798, MAMMA1000842, MAMMA1000859,

2812

MAMMA1000897, MAMMA1000956, MAMMA1001008, MAMMA1001030, MAMMA1001041, MAMMA1001073, MAMMA1001080, MAMMA1001139, MAMMA1001154, MAMMA1001322, MAMMA1001388, MAMMA1001411, MAMMA1001487, MAMMA1001751, MAMMA1001754, MAMMA1001771, MAMMA1002009, MAMMA1002427, MAMMA1002428, MAMMA1002461, MAMMA1002524, MAMMA1002573, MAMMA1002598, MAMMA1002655, MAMMA1002684, MAMMA1002769, MAMMA1002844, MAMMA1002881, MAMMA1002890, MAMMA1002938, MAMMA1002947, MAMMA1003035, MAMMA1003089, MAMMA1003146, MAMMA1003150, NT2RM1000035, NT2RM1000037, NT2RM1000062, NT2RM1000080, NT2RM1000092, NT2RM1000131, NT2RM1000199, NT2RM1000257, NT2RM1000260, NT2RM1000355, NT2RM1000430, NT2RM1000563, NT2RM1000648, NT2RM1000742, NT2RM1000770, NT2RM1000800, NT2RM1000811, NT2RM1000833, NT2RM1000857, NT2RM1000867, NT2RM1000882, NT2RM1000905, NT2RM1001008,

【1053】NT2RM1001115, NT2RM1001139, NT2RM2000259, NT2RM2000260, NT2RM2000287, NT2RM2000395, NT2RM2000402, NT2RM2000407, NT2RM2000422, NT2RM2000490, NT2RM2000522, NT2RM2000566, NT2RM2000581, NT2RM2000609, NT2RM2000821, NT2RM2001370, NT2RM2001393, NT2RM2001499, NT2RM2001547, NT2RM2001613, NT2RM2001648, NT2RM2001659, NT2RM2001671, NT2RM2001688, NT2RM2001698, NT2RM2001718, NT2RM2001753, NT2RM2001760, NT2RM2001785, NT2RM2001930, NT2RM2001950, NT2RM2001997, NT2RM2001998, NT2RM2002049, NT2RM2002145, NT2RM4000233, NT2RM4000433, NT2RM4000457, NT2RM4000486, NT2RM4000496, NT2RM4000520, NT2RM4000634, NT2RM4000674, NT2RM4000700, NT2RM4000764, NT2RM4000778, NT2RM4000795, NT2RM4000820, NT2RM4000857, NT2RM4001032, NT2RM4001054, NT2RM4001116, NT2RM4001455, NT2RM4001666, NT2RM4001810, NT2RM4001813, NT2RM4001930, NT2RM4001987, NT2RM4002054, NT2RM4002073, NT2RM4002145, NT2RM4002146, NT2RM4002189, NT2RM4002194, NT2RM4002251, NT2RM4002339, NT2RM4002438, NT2RM4002446, NT2RM4002452, NT2RM4002460, NT2RM4002493, NT2RM4002558, NT2RM4002565, NT2RM4002571, NT2RM4002594, NT2RP1000130, NT2RP1000191, NT2RP100026, NT2RP1000358, NT2RP1000413, NT2RP1000418, NT2RP1000547, NT2RP1000609, NT2RP1000677, NT2RP1000767, NT2RP1000782, NT2RP1000856, NT2RP1001113, NT2RP1001247, NT2RP1001286, NT2RP1001310, NT2RP1001311, NT2RP1001313, NT2RP1001385, NT2RP1001449, NT2RP1001546, NT2RP1001569, NT2RP2000032, NT2RP2000040, NT2RP2000056,

【1054】NT2RP2000070, NT2RP2000091, NT2RP2000114, NT2RP2000120, NT2RP2000173, NT2RP2000175, NT2RP2000195, NT2RP2000257, NT2RP2000270, NT2RP2000283, NT2RP2000288, NT2RP2000289, NT2RP2000459, NT2RP

2813

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2814

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2815

6, PLACE1007484, PLACE1007583, PLACE1007632, PLACE1007645, PLACE1007649, PLACE1007852, PLACE1007877, PLACE1007954, PLACE1008273, PLACE1008309, PLACE1008331, PLACE1008402, PLACE1008424, PLACE1008429, PLACE1008531, PLACE1008532, PLACE1008533, PLACE1008568, PLACE1008643, PLACE1008693, PLACE1008715, PLACE1009045, PLACE1009094, PLACE1009298, PLACE1009319, PLACE1009338, PLACE1009368, PLACE1009493, PLACE1009639, PLACE1009659, PLACE1009708, PLACE1009731, PLACE1009845, PLACE1009861, PLACE1009935, PLACE1009992, PLACE1010089, PLACE1010231, PLACE1010321, PLACE1010362,

【1058】PLACE1010599, PLACE1010622, PLACE1010662, PLACE1010811, PLACE1010917, PLACE1010942, PLACE1010954, PLACE1011090, PLACE1011214, PLACE1011221, PLACE1011371, PLACE1011399, PLACE1011492, PLACE1011646, PLACE1011749, PLACE1011896, PLACE2000034, PLACE2000062, PLACE2000111, PLACE2000132, PLACE2000176, PLACE2000187, PLACE2000216, PLACE2000335, PLACE2000341, PLACE2000373, PLACE2000379, PLACE2000398, PLACE2000399, PLACE2000425, PLACE2000438, PLACE2000458, PLACE2000477, PLACE3000020, PLACE3000021, PLACE3000226, PLACE3000242, PLACE3000244, PLACE3000339, PLACE3000373, PLACE3000399, PLACE3000406, PLACE3000413, PLACE3000455, PLACE4000052, PLACE4000063, PLACE4000129, PLACE4000247, PLACE4000250, PLACE4000259, PLACE4000300, PLACE4000387, PLACE4000431, PLACE4000487, PLACE4000494, PLACE4000522, PLACE4000548, PLACE4000581, PLACE4000593, PLACE4000650, THYR01000156, THYR01000327, THYR01000394, THYR01000395, THYR01000570, THYR01000748, THYR01000756, THYR01000783, THYR01001134, THYR01001271, THYR01001287, THYR01001320, THYR01001401, THYR01001534, THYR01001537, THYR01001541, THYR01001828, Y79AA1000258, Y79AA1000420, Y79AA1000469, Y79AA1000734, Y79AA1000800, Y79AA1000976, Y79AA1001023, Y79AA1001177, Y79AA1001384, Y79AA1001394, Y79AA1001603, Y79AA1001647, Y79AA1001846, Y79AA1001874, Y79AA1002139, Y79AA1002246, Y79AA1002351, Y79AA1002399, Y79AA1002416,

【1059】糖タンパク質関連タンパク質に分類されたクローンは以下の141クローンであった。

HEMBA1000156, HEMBA1000518, HEMBA1000852, HEMBA1001071, HEMBA1001286, HEMBA1001661, HEMBA1001734, HEMBA1001866, HEMBA1003071, HEMBA1003077, HEMBA1003281, HEMBA1003538, HEMBA1003679, HEMBA1003866, HEMBA1005576, HEMBA1005581, HEMBA1005699, HEMBA1006038, HEMBA1006976, HEMBA1007301, HEMBB1000317, HEMBB1000915, HEMBB1001871, HEMBB1001872, HEMBB1002193, MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA10

2816

01388, MAMMA1002329, MAMMA1002428, MAMMA1002573, MAMMA1003150, NT2RM1000648, NT2RM1001115, NT2RM2000260, NT2RM2000407, NT2RM2000422, NT2RM2000490, NT2RM2001499, NT2RM2001659, NT2RM2001930, NT2RM40000820, NT2RM4000857, NT2RM4001810, NT2RM4001813, NT2RM4001987, NT2RM4002145, NT2RM4002189, NT2RM4002251, NT2RM4002460, NT2RM4002558, NT2RP1000677, NT2RP1000782, NT2RP1000856, NT2RP1001546, NT2RP2000056, NT2RP2000070, NT2RP2001295, NT2RP2001378, NT2RP2001597, NT2RP2001991, NT2RP2002025, NT2RP2002078, NT2RP2002385, NT2RP2004587, NT2RP2004732, NT2RP2005531, NT2RP3000207, NT2RP3000531, NT2RP3000825, NT2RP3001140, NT2RP3002810, NT2RP3003672, NT2RP3003701, NT2RP3003716, NT2RP3003914, NT2RP3004148, NT2RP4000212, NT2RP4000417, NT2RP4000724, NT2RP4000817, NT2RP4000925, NT2RP4001150, NT2RP4001372, NT2RP4001730, NT2RP4001822, NT2RP4001823, NT2RP5003522, OVARC1000091, OVARC1000288, OVARC1000682, OVARC1001055, OVARC1001506, OVARC1001713, OVARC1002127, PLACE1000213, PLACE1000401, PLACE1002437, PLACE1002583, 【1060】PLACE1002722, PLACE1003045, PLACE1003238, PLACE1003258, PLACE1003493, PLACE1004197, PLACE1004793, PLACE1005953, PLACE1005955, PLACE1006157, PLACE1006239, PLACE1006368, PLACE1006534, PLACE1006754, PLACE1006956, PLACE1007416, PLACE1007632, PLACE1007649, PLACE1008643, PLACE1009094, PLACE1009992, PLACE1010231, PLACE1010662, PLACE1011371, PLACE2000034, PLACE2000373, PLACE2000398, PLACE2000399, PLACE2000438, PLACE2000458, PLACE3000339, PLACE4000063, PLACE4000230, PLACE4000522, PLACE4000548, PLACE4000581, THYR01000327, THYR01000756, THYR01001287, Y79AA1001603, Y79AA1001874

【1061】シグナル伝達関連タンパク質に分類されたクローンは以下の129クローンであった。

HEMBA1000303, HEMBA1000369, HEMBA1000608, HEMBA1000657, HEMBA1000919, HEMBA1001019, HEMBA1001174, HEMBA1001822, HEMBA1001921, HEMBA1002139, HEMBA1002212, HEMBA1002341, HEMBA1002417, HEMBA1002768, HEMBA1003250, HEMBA1003291, HEMBA1003645, HEMBA1004286, HEMBA1005737, HEMBA1006130, HEMBA1006708, HEMBB1000083, HEMBB1000266, HEMBB1000632, HEMBB1000781, HEMBB1000831, HEMBB1002193, MAMMA1000173, MAMMA1001038, MAMMA1001198, MAMMA1002842, MAMMA1003057, NT2RM1000702, NT2RM1000772, NT2RM1001072, NT2RM2000030, NT2RM2000469, NT2RM2000612, NT2RM2001221, NT2RM2001345, NT2RM2002128, NT2RM4000229, NT2RM4000354, NT2RM4000611, NT2RM4000798, NT2RM4001411, NT2RM4001412, NT2RM4001629, NT2RM4001758, NT2RM4002013, NT2RM4002527, NT2RP1000018, NT2RP1000701, NT2RP1001294, NT2RP1001302, NT2RP2000668, NT2RP2001440, N

2817

T2RP2001560, NT2RP2002058, NT2RP2002193, NT2RP2002408, NT2RP2002710, NT2RP2002929, NT2RP2003164, NT2RP2003912, NT2RP2004232, NT2RP2004768, NT2RP2006071, NT2RP2006534, NT2RP3000759, NT2RP3000845, NT2RP3001646, NT2RP3001857, NT2RP3001938, NT2RP3002004, NT2RP3002785, NT2RP3002909, NT2RP3002988, NT2RP3003800, NT2RP3004189, NT2RP3004544, NT2RP4000147, NT2RP4000839, NT2RP4001122, NT2RP4001148, NT2RP4001336, NT2RP4001375, NT2RP4001644, NT2RP4001725, NT2RP4001849, NT2RP4001896, NT2RP4001927, NT2RP4002408, NT2RP5003477, OVARC1000013, OVARC1000437, OVARC1000556, OVARC1000649, OVARC1000945, OVARC1001200, 【1062】OVARC1002182, PLACE1000977, PLACE1001387, PLACE1002493, PLACE1002591, PLACE1003190, PLACE1003353, PLACE1004128, PLACE1004302, PLACE1004937, PLACE1005243, PLACE1008000, PLACE1008244, PLACE1008650, PLACE1009468, PLACE1009596, PLACE1009708, PLACE1009845, PLACE1010926, PLACE1011041, PLACE200164, PLACE2000371, PLACE3000145, PLACE3000350, THYR01000072, THYR01000748, THYR01001120, Y79AA1000328, Y79AA1002431

【1063】転写関連タンパク質に分類されたクローンは以下の309クローンであった。

HEMBA1000158, HEMBA1000201, HEMBA1000216, HEMBA1000555, HEMBA1000561, HEMBA1000851, HEMBA1001077, HEMBA1001137, HEMBA1001405, HEMBA1001510, HEMBA1001635, HEMBA1001804, HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1001869, HEMBA1002035, HEMBA1002092, HEMBA1002177, HEMBA1002770, HEMBA1002935, HEMBA1003408, HEMBA1003545, HEMBA1003568, HEMBA1003662, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004353, HEMBA1004389, HEMBA1004479, HEMBA1004758, HEMBA1004973, HEMBA1005219, HEMBA1005359, HEMBA1005513, HEMBA1005528, HEMBA1005548, HEMBA1005558, HEMBA1005931, HEMBA1006158, HEMBA1006248, HEMBA1006278, HEMBA1006283, HEMBA1006347, HEMBA1006359, HEMBA1006559, HEMBA1006941, HEMBB1000789, HEMBB1001011, HEMBB1001314, HEMBB1001482, HEMBB1001673, HEMBB1001749, HEMBB1001839, HEMBB1001908, HEMBB1002134, HEMBB1002217, HEMBB1002342, HEMBB1002607, MAMMA1000183, MAMMA1000388, MAMMA1001105, MAMMA1001222, MAMMA1001260, MAMMA1001627, MAMMA1001633, MAMMA1001743, MAMMA1001820, MAMMA1001837, MAMMA1002617, MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000086, NT2RM1000746, NT2RM1000885, NT2RM1000894, NT2RM1001092, NT2RM2000013, NT2RM2000452, NT2RM2000735, NT2RM2000740, NT2RM2001035, NT2RM2001105, NT2RM2001575, NT2RM2001670, NT2RM2001716, NT2RM2001771, NT2RM2002091, NT2RM40000024, NT2RM4000046, NT2RM4000104, NT2RM4000202, NT2RM

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4000531, NT2RM4000595, NT2RM4000733, NT2RM4000734, 【1064】NT2RM4000741, NT2RM4000751, NT2RM4000996, NT2RM4001092, NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RM4002066, NT2RP1000086, NT2RP1000111, NT2RP1000574, NT2RP1000902, NT2RP1001013, NT2RP2000008, NT2RP200126, NT2RP2000297, NT2RP2000420, NT2RP2001174, NT2RP2001233, NT2RP2001756, NT2RP2001869, NT2RP2002046, NT2RP2002252, NT2RP2002270, NT2RP2002464, NT2RP2002503, NT2RP2002520, NT2RP2002591, NT2RP2002880, NT2RP2002939, NT2RP2002993, NT2RP2003243, NT2RP2003329, NT2RP2003347, NT2RP2003480, NT2RP2003522, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004066, NT2RP2004187, NT2RP2004920, NT2RP2004961, NT2RP2005003, NT2RP2005139, NT2RP2005325, NT2RP2005496, NT2RP2005701, NT2RP2005722, NT2RP2005776, NT2RP2005942, NT2RP2006238, NT2RP2006436, NT2RP30000050, NT2RP3000320, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP3000603, NT2RP3000605, NT2RP3000632, NT2RP3001057, NT2RP3001107, NT2RP3001111, NT2RP3001120, NT2RP3001150, NT2RP3001268, NT2RP3001338, NT2RP3001398, NT2RP3001527, NT2RP3001688, NT2RP3001855, NT2RP3002165, NT2RP3002399, NT2RP3002876, NT2RP3003133, NT2RP3003193, NT2RP3003251, NT2RP3003331, NT2RP3003327, NT2RP3003555, NT2RP3004016, NT2RP3004125, NT2RP3004242, NT2RP3004428, NT2RP3004498, NT2RP3004566, NT2RP3004617, NT2RP4000210, NT2RP4000398, NT2RP4000455, NT2RP4000648, NT2RP4000837, NT2RP4000865.

【1065】NT2RP4000997, NT2RP4001029, NT2RP4001080, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2RP4001551, NT2RP4001568, NT2RP4001638, NT2RP4001753, NT2RP4001760, NT2RP4001790, NT2RP4001838, NT2RP4001938, NT2RP4002078, NT2RP4002081, NT2RP5003461, OVARC1000151, OVARC1000241, OVARC1000479, OVARC1001271, OVARC1001417, OVARC1001436, PLACE1000133, PLACE1000583, PLACE1000706, PLACE1000786, PLACE1000979, PLACE1001118, PLACE1001238, PLACE1001294, PLACE1001304, PLACE1001383, PLACE1001602, PLACE1001632, PLACE1002171, PLACE1002438, PLACE1002450, PLACE1002532, PLACE1002775, PLACE1002834, PLACE1003302, PLACE1003605, PLACE1003738, PLACE1003885, PLACE1004471, PLACE1005584, PLACE1005803, PLACE1005966, PLACE1006167, PLACE1006318, PLACE1006438, PLACE1006482, PLACE1007239, PLACE1007346, PLACE1007488, PLACE1007547, PLACE1007598, PLACE1007955, PLACE1008132, PLACE1008201, PLACE1009099, PLACE1009246, PLACE1009308, PLACE1009398, PLACE1009798, PLACE1010134, PLACE1010702, PLACE1010771, PLACE1010870, PLACE10

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11160, PLACE1011433, PLACE1011576, PLACE3000009, PLACE3000169, PLACE3000254, PLACE4000128, PLACE4000156, PLACE4000192, PLACE4000211, PLACE4000261, PLACE4000450, PLACE4000489, THYRO1000085, THYRO1000121, THYRO1000242, THYRO1000488, THYRO1000501, THYRO1000569, THYRO1001100, THYRO1001189, THYRO1001809, Y79AA1000013, Y79AA1000033, Y79AA1000037, Y79AA1000342, Y79AA1000627, Y79AA1000705, Y79AA1001299, Y79AA1001312, Y79AA1001391, Y79AA1001533, Y79AA1001613, Y79AA1001866, Y79AA1002103, Y79AA1002229, Y79AA1002433, Y79AA1002472, Y79AA1002482,

【1066】疾患関連タンパク質に分類されたクローンは以下の392クローンであった。

HEMBA1000020, HEMBA1000216, HEMBA1000304, HEMBA1000561, HEMBA1000569, HEMBA1000910, HEMBA1001043, HEMBA1001059, HEMBA1001071, HEMBA1001088, HEMBA1001569, HEMBA1001661, HEMBA1001672, HEMBA1001819, HEMBA1001921, HEMBA1002267, HEMBA1002419, HEMBA1002469, HEMBA1002547, HEMBA1002555, HEMBA1002810, HEMBA1002939, HEMBA1002997, HEMBA1003148, HEMBA1003369, HEMBA1003417, HEMBA1003418, HEMBA1003433, HEMBA1003538, HEMBA1003555, HEMBA1003568, HEMBA1003569, HEMBA1003581, HEMBA1004168, HEMBA1004202, HEMBA1004248, HEMBA1004275, HEMBA1004321, HEMBA1004353, HEMBA1004356, HEMBA1004479, HEMBA1004509, HEMBA1004669, HEMBA1005009, HEMBA1005338, HEMBA1005367, HEMBA1005423, HEMBA1005528, HEMBA1005581, HEMBA1005621, HEMBA1005699, HEMBA1006507, HEMBA1006650, HEMBA1006652, HEMBA1006737, HEMBA1006807, HEMBA1006877, HEMBA1007121, HEMBA1007243, HEMBB1000119, HEMBB1000693, HEMBB1000927, HEMBB1000985, HEMBB1001068, HEMBB1001282, HEMBB1001339, HEMBB1001482, HEMBB1001564, HEMBB1001802, HEMBB1001905, HEMBB1001908, HEMBB1002217, HEMBB1002477, MAMMA1000388, MAMMA1000731, MAMMA1001305, MAMMA1001633, MAMMA1001868, MAMMA1002170, MAMMA1002198, MAMMA1002268, MAMMA1002485, MAMMA1002530, MAMMA1002858, MAMMA1002869, MAMMA1002881, MAMMA1003047, MAMMA1003146, MAMMA1003166, NT2RM1000001, NT2RM1000153, NT2RM1000252, NT2RM1000555, NT2RM1000770, NT2RM1000826, NT2RM1000850, NT2RM1001003, NT2RM1001092, NT2RM1001102, NT2RM2000191,

【1067】NT2RM2000363, NT2RM2000594, NT2RM2000624, NT2RM2000714, NT2RM2000821, NT2RM2001035, NT2RM2001575, NT2RM2001652, NT2RM2001664, NT2RM2001668, NT2RM2001698, NT2RM2001803, NT2RM2001839, NT2RM4000155, NT2RM4000471, NT2RM4000486, NT2RM4000657, NT2RM4000751, NT2RM4000996, NT2RM4001629, NT2RM4001810, NT2RM4001819, NT2RM4001865, NT2RM4001876, NT2RM4001940, NT2RM4002066, NT2RM4002093, NT2RM4002146, NT2RM4002161, NT2RM4002323, NT2RM4002558, NT

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RM4002571, NT2RP1000086, NT2RP1000574, NT2RP1000738, NT2RP1000825, NT2RP1000833, NT2RP1000959, NT2RP1000966, NT2RP1001013, NT2RP1001185, NT2RP1001482, NT2RP1001665, NT2RP2000070, NT2RP2000147, NT2RP2000224, NT2RP2000248, NT2RP2000297, NT2RP2000310, NT2RP2000414, NT2RP2000420, NT2RP2000523, NT2RP2000809, NT2RP2000812, NT2RP2001233, NT2RP2001327, NT2RP2001378, NT2RP2001394, NT2RP2001397, NT2RP2001460, NT2RP2001520, NT2RP2001536, NT2RP2001876, NT2RP2001898, NT2RP2002025, NT2RP2002058, NT2RP2002124, NT2RP2002325, NT2RP2002503, NT2RP2002959, NT2RP2003000, NT2RP2003157, NT2RP2003164, NT2RP2003228, NT2RP2003295, NT2RP2003517, NT2RP2003564, NT2RP2003604, NT2RP2003714, NT2RP2003737, NT2RP2003952, NT2RP2004013, NT2RP2004170, NT2RP2004587, NT2RP2004732, NT2RP2004933, NT2RP2005003, NT2RP2005144, NT2RP2005239, NT2RP2005276, NT2RP2005288, NT2RP2005315, NT2RP2005325, NT2RP2005336, NT2RP2005358, NT2RP2005407, NT2RP2005436, NT2RP2005476, NT2RP2005525, NT2RP2005694,

【1068】NT2RP2005719, NT2RP2006043, NT2RP2006071, NT2RP2006219, NT2RP2006312, NT2RP2006456, NT2RP3000050, NT2RP3000068, NT2RP3000085, NT2RP3000299, NT2RP3000403, NT2RP3000596, NT2RP3000739, NT2RP3000753, NT2RP3000875, NT2RP3001057, NT2RP3001081, NT2RP3001216, NT2RP3001307, NT2RP3001338, NT2RP3001427, NT2RP3001428, NT2RP3001679, NT2RP3001723, NT2RP3001855, NT2RP3001898, NT2RP3001969, NT2RP3002056, NT2RP3002062, NT2RP3002151, NT2RP3002351, NT2RP3002399, NT2RP3002953, NT2RP3002988, NT2RP3003078, NT2RP3003251, NT2RP3003282, NT2RP3003313, NT2RP3003327, NT2RP3003409, NT2RP3003672, NT2RP3003831, NT2RP3004016, NT2RP3004078, NT2RP3004209, NT2RP3004258, NT2RP3004490, NT2RP3004534, NT2RP3004569, NT2RP3004572, NT2RP4000109, NT2RP4000367, NT2RP4000376, NT2RP4000449, NT2RP4000855, NT2RP4000879, NT2RP4000925, NT2RP4001086, NT2RP4001126, NT2RP4001150, NT2RP4001213, NT2RP4001276, NT2RP4001407, NT2RP4001433, NT2RP4001483, NT2RP4001575, NT2RP4001760, NT2RP4001861, NT2RP4002078, NT2RP4002791, OVARC1000014, OVARC1000139, OVARC1000520, OVARC1000722, OVARC1000771, OVARC1000834, OVARC1001051, OVARC1001113, OVARC1001244, OVARC1001372, OVARC1001417, OVARC1001496, OVARC1001506, OVARC1001577, OVARC1001726, OVARC1001766, OVARC1001809, OVARC1002165, PLACE1000133, PLACE1000383, PLACE1000420, PLACE1000583, PLACE1000588, PLACE1001171, PLACE1001387, PLACE1001602, PLACE1002046, PLACE1002140, PLACE1002437, PLACE1002474,

【1069】PLACE1002685, PLACE1002782, PLACE10028

2821

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2822

NT2RM2001035(604913), NT2RM2001575(109092), NT2RM2001652(604141), NT2RM2001664(603722), NT2RM2001668(602952), NT2RM2001698(604327), NT2RM2001803(603722), NT2RM2001839(603420), NT2RM4000155(187790), NT2RM40000471(603485), NT2RM4000486(168730), NT2RM4000657(602142), NT2RM4000751(602277), NT2RM4000996(603971), NT2RM4001629(601114), NT2RM4001810(155760), NT2RM4001819(176873), NT2RM4001876(179555), NT2RM4001940(603887), NT2RM4002066(300188), NT2RM4002093(600693), NT2RM4002146(602603), NT2RM4002161(254780), NT2RM4002558(604194), NT2RM4002571(602274), NT2RP1000086(602219), NT2RP1000574(601740), NT2RP1000825(602732), NT2RP1000833(602973), NT2RP1000959(180510), NT2RP1000966(164035), NT2RP1001013(194558), NT2RP1001185(243500), NT2RP1001482(600586), NT2RP1001665(114180), NT2RP2000070(600976), NT2RP2000147(603535), NT2RP2000248(603367), NT2RP2000297(602277), NT2RP2000310(239500), NT2RP2000414(601037), NT2RP2000420(600834), NT2RP2000523(600130), NT2RP2000809(603885), NT2RP2000812(160777), NT2RP2001233(603971), NT2RP2001327(191161), NT2RP2001378(158370), NT2RP2001394(300208), NT2RP2001397(602755), NT2RP2001460(190370), NT2RP2001520(603667), NT2RP2001536(600675), NT2RP2001876(601833), NT2RP2001898(147264), NT2RP2002025(601581), NT2RP2002058(604737), 【1071】NT2RP2002124(603486), NT2RP2002325(603866), NT2RP2002503(601781), NT2RP2002959(602962), NT2RP2003000(191161), NT2RP2003157(601940), NT2RP2003164(604746), NT2RP2003228(602638), NT2RP2003295(603494), NT2RP2003517(190040), NT2RP2003564(109092), NT2RP2003604(604785), NT2RP2003714(603971), NT2RP2003737(602962), NT2RP2003952(602675), NT2RP2004013(602542), NT2RP2004170(300196), NT2RP2004587(162250), NT2RP2004732(162250), NT2RP2004933(603289), NT2RP2005003(109092), NT2RP2005144(604730), NT2RP2005239(603485), NT2RP2005276(602371), NT2RP2005288(603524), NT2RP2005315(604039), NT2RP2005325(603759), NT2RP2005336(190370), NT2RP2005358(603573), NT2RP2005407(167040), NT2RP2005436(601940), NT2RP2005476(602680), NT2RP2005525(602655), NT2RP2005719(601178), NT2RP2006043(601940), NT2RP2006071(604299), NT2RP2006219(601279), NT2RP2006312(603111), NT2RP2006456(604619), NT2RP3000050(603971), NT2RP3000068(182530), NT2RP3000085(300032), NT2RP3000029(602941), NT2RP3000403(604981), NT2RP3000596(190370), NT2RP3000739(125370), NT2RP3000753(162230), NT2RP3001057(603971), NT2RP3001081(603524), NT2RP3001216(603121), NT2RP3001307(180069), NT2RP3001338(314998), NT2RP3001428(189940), NT2RP3001723(60456

2823

9), NT2RP3001855(602100), NT2RP3001898(604561), NT2RP3001969(190370), NT2RP3002056(180201), NT2RP3002062(603885), NT2RP3002151(139259), NT2RP3002351(604887), NT2RP3002399(602638), NT2RP3002953(604967), NT2RP3002988(603258), NT2RP3003251(109092), NT2RP3003282(602378), NT2RP3003313(603810), NT2RP3003409(604533), NT2RP3003672(313470), NT2RP3003831(604051), NT2RP3004016(601742), NT2RP3004078(142765), NT2RP3004209(125255), NT2RP3004258(604347), NT2RP3004490(603328), NT2RP3004534(600586), NT2RP3004569(106410), NT2RP3004572(604912), NT2RP4000109(603745), NT2RP4000367(603722), NT2RP4000376(603873), NT2RP4000449(604479), NT2RP4000855(602675), NT2RP4000879(314370), NT2RP4000925(600245), NT2RP4001086(162230), NT2RP4001126(190370), NT2RP4001150(601581), NT2RP4001213(602277), NT2RP4001276(190370), NT2RP4001407(190370), NT2RP4001433(602277), NT2RP4001483(203740), NT2RP4001575(603443), NT2RP4001760(305400), NT2RP4001861(190370), NT2RP4002078(603971), NT2RP4002791(189940), OVARC1000014(603371), OVARC1000139(603486),

【1072】OVARC1000520(604126), OVARC1000722(604014), OVARC1000771(179509), OVARC1001051(600051), OVARC1001113(602121), OVARC1001244(601540), OVARC1001372(603145), OVARC1001417(300182), OVARC1001496(602619), OVARC1001506(601313), OVARC1001577(603269), OVARC1001726(300103), OVARC1001766(603910), OVARC1001809(603730), PLACE1000133(602542), PLACE1000383(300171), PLACE1000420(600312), PLACE1000583(194558), PLACE1000588(600411), PLACE1001171(310400), PLACE1001387(600206), PLACE1001602(604913), PLACE1002046(151625), PLACE1002140(603748), PLACE1002437(600046), PLACE1002474(602108), PLACE1002685(604515), PLACE1002782(602095), PLACE1002834(194558), PLACE1002908(604327), PLACE1003045(173910), PLACE1003302(194558), PLACE1003353(604704), PLACE1003366(603681), PLACE1003493(601456), PLACE1003669(190370), PLACE1003704(601940), PLACE1003903(123860), PLACE1003968(602742), PLACE1004183(604701), PLACE1004197(601610), PLACE1004277(603493), PLACE1004316(604261), PLACE1004358(603272), PLACE1004471(194558), PLACE1004506(603450), PLACE1004510(604912), PLACE1004674(601057), PLACE1004777(118423), PLACE1004814(601940), PLACE1005494(603652), PLACE1006040(603061), PLACE1006170(601026), PLACE1006438(600834), PLACE1006615(603910), PLACE1007140(190370), PLACE1007239(604784), PLACE1007257(300108), PLACE1007511(148020), PLACE1007598(602277), PLACE1008177(190370), PLACE1008356(604039), PLACE1008402(603344), PLACE1008696(602141), PLACE1009027(300121), PL

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【1073】THYR01001458(160776), THYR01001617(602744), THYR01001671(603281), Y79AA1000346(604355), Y79AA1000469(602434), Y79AA1000560(601026), Y79AA1000734(603867), Y79AA1000782(600417), Y79AA1001391(142959), Y79AA1001548(600286), Y79AA1001594(600936), Y79AA1001711(600063), Y79AA1001874(600315), Y79AA1002204(605033), Y79AA1002210(191161), Y79AA1002472(603971), Y79AA1002482(603971),

【1074】酵素・代謝関連タンパク質のカテゴリーに属すると推定されたクローンは以下の425クローンであった。

HEMBA1000012, HEMBA1000129, HEMBA1000141, HEMBA1000150, HEMBA1000542, HEMBA1000852, HEMBA1001019, HEMBA1001257, HEMBA1001526, HEMBA1001620, HEMBA1001866, HEMBA1001896, HEMBA1002212, HEMBA1002513, HEMBA1002746, HEMBA1002973, HEMBA1003046, HEMBA1003136, HEMBA1003179, HEMBA1003250, HEMBA1003291, HEMBA1003408, HEMBA1003538, HEMBA1003679, HEMBA1003680, HEMBA1004199, HEMBA1004227, HEMBA1004408, HEMBA1004509, HEMBA1004734, HEMBA1004768, HEMBA1005394, HEMBA1005513, HEMBA1005737, HEMBA1005815, HEMBA1006031, HEMBA1006272, HEMBA1006278, HEMBA1006291, HEMBA1006309, HEMBA1006347, HEMBA1006485, HEMBA1006521, HEMBA1006624, HEMBA1006885, HEMBA1006976, HEMBA1007121, HEMBA1007224, HEMBA1007243, HEMBA1007300, HEMBB1000083, HEMBB1000217, HEMBB1000915, HEMBB1000947, HEMBB1001137, HEMBB1001346, HEMBB1001429, HEMBB1001443, HEMBB1001915, HEMBB1001950, HEMBB1002042, MAMMA1000020, MAMMA1000085, MAMMA1000672, MAMMA1000713, MAMMA1000841, MAMMA1000897, MAMMA1001008, MAMMA1001038, MAMMA1001059, MAMMA1001476, MAMMA1001501, MAMMA1002268, MAMMA1002470, MAMMA1002530, MAMMA1002573, MAMMA1002619, MAMMA1002655, MAMMA1002671, MAMMA1003013, MAMMA1003035, NT2RM1000039, NT

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2826

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【1079】ATP・GTP結合タンパク質のカテゴリに属すると推定されたクローンは、以下の217クローンであった。

HEMBA1000012, HEMBA1000129, HEMBA1000185, HEMBA1000491, HEMBA1000531, HEMBA1001019, HEMBA1001174, HEMBA1001387, HEMBA1001595, HEMBA1001723, HEMBA1001913, HEMBA1002161, HEMBA1002212, HEMBA1002876, HEMBA1002997, HEMBA1003250, HEMBA1003291, HEMBA1003369, HEMBA1003555, HEMBA1003560, HEMBA1004131, HEMBA1004199, HEMBA1004202, HEMBA1004354, HEMBA1004697, HEMBA1005047, HEMBA1005595, HEMBA1007018, HEMBA1007151, HEMBB1000083, HEMBB1000226, HEMBB1000264, HEMBB1000632, HEMBB1000725, HEMBB1001294, HEMBB1002193, MAMMA1000085, MAMMA1000612, MAMMA1000731, MAMMA1000738, MAMMA1001038, MAMMA1001735, MAMMA1001768, MAMMA1003127, NT2RM1000187, NT2RM1000388, NT2RM1000702, NT2RM1000772, NT2RM1000924, NT2RM2000469, NT2RM2000577, NT2RM2000740, NT2RM2001100, NT2RM2001201, NT2RM2001345, NT2RM2001823, NT2RM2002128, NT2RM4000155, NT2RM4000191, NT2RM4000356, NT2RM4000496, NT2RM4000611, NT2RM4000733, NT2RM4000820, NT2RM4001084, NT2RM4001178, NT2RM4001344, NT2RM4001444, NT2RM4001592, NT2RM4001714, NT2RM4001758, NT2RM4001880, NT2RM4002062, NT2RM4002174, NT2RM4002205, NT2RM4002527, NT2RM4002594, NT2RM4002623, NT2RP1000470, NT2RP1000478, NT2RP1000915, NT2RP1000958, NT2RP1001080, NT2RP1001410, NT2RP1001569, NT2RP2000126, NT2RP2000258, NT2RP2000329, NT2RP2000660, NT2RP2000668, NT2RP2000710, NT2RP2000812, NT2RP2000880, NT2RP2001245, NT2RP2001392, NT2RP2002606, NT2RP2003277, NT2RP2003912, NT2RP2004538, NT2RP2004568, 【1080】NT2RP2004689, NT2RP2004768, NT2RP2004791, NT2RP2004920, NT2RP2005344, NT2RP2005393, NT2RP2005763, NT2RP2006534, NT2RP3000046, NT2RP3000025, NT2RP3000350, NT2RP3000359, NT2RP3000366, NT2RP3000397, NT2RP3000759, NT2RP3000845, NT2RP3000875, NT2RP3001150, NT2RP3001427, NT2RP3001453, NT2RP3001529, NT2RP3001730, NT2RP3001799, NT2RP3001857, NT2RP3001938, NT2RP3002007, NT2RP3002151, NT2RP3002330, NT2RP3002399, NT2RP3002671, NT2RP3003301, NT2RP3003353, NT2RP3003589, NT2RP3003809, NT2RP3003876, NT2RP3004189, NT2RP3004428, NT2RP3004578, NT2RP4000290, NT2RP4000481, NT2RP4000518, NT2RP4000781, NT2RP4000839, NT2RP4000929, NT2RP4001041, NT2RP4001079, NT2RP4001375, NT2RP4001414, NT2RP4001592, NT2RP4001634, NT2RP4001644, NT2RP4001656, NT2RP4001896, NT2RP4002047, NT2RP4002058, NT2RP4002408, NT2RP5003477, OVARC1000013, OVARC1000304, OVARC1000556, OVARC1000771, OVARC1000800, OVARC1001068, OVARC1002138, PLACE1000040, PLACE1000588, PLACE1001104, 50

PLACE1001739, PLACE1002433, PLACE1002437, PLACE1002714, PLACE1003394, PLACE1003521, PLACE1003915, PLACE1004902, PLACE1005243, PLACE1005305, PLACE1005549, PLACE1005739, PLACE1005921, PLACE1006119, PLACE1006196, PLACE1006552, PLACE1006956, PLACE1007409, PLACE1007697, PLACE1007946, PLACE1008244, PLACE1009404, PLACE1009476, PLACE1009596, PLACE1009908, PLACE1010134, PLACE1010720, PLACE1010896, PLACE1011109, PLACE1011114, PLACE1011310, PLACE1011922, PLACE2000014, PLACE2000039, PLACE2000274, PLACE2000404, PLACE2000427, PLACE3000350, PLACE4000009, PLACE4000014, PLACE4000326, SKNMC1000013, THYR01000072, THYR01001458, Y79AA1000833, Y79AA1000962, Y79AA1001394, Y79AA1001875, Y79AA1001963, Y79AA1002209, 【1081】核タンパク質のカテゴリに属すると推定されたクローンは、以下の320クローンであった。

HEMBA1000005, HEMBA1000158, HEMBA1000216, HEMBA1000561, HEMBA1000591, HEMBA1001088, HEMBA1001137, HEMBA1001405, HEMBA1001510, HEMBA1001579, HEMBA1001809, HEMBA1001819, HEMBA1001824, HEMBA1001847, HEMBA1001869, HEMBA1002177, HEMBA1002241, HEMBA1002495, HEMBA1002569, HEMBA1002935, HEMBA1002951, HEMBA1002999, HEMBA1003408, HEMBA1003545, HEMBA1003662, HEMBA1003684, HEMBA1003690, HEMBA1003760, HEMBA1004203, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA1004973, HEMBA1005219, HEMBA1005359, HEMBA1005558, HEMBA1005931, HEMBA1006278, HEMBA1006283, HEMBA1006359, HEMBA1006485, HEMBA1007087, HEMBB1000226, HEMBB1000789, HEMBB1001011, HEMBB1001056, HEMBB1001242, HEMBB1001482, HEMBB1001915, HEMBB1002134, HEMBB1002217, MAMMA1000183, MAMMA1000731, MAMMA1001105, MAMMA1001222, MAMMA1001260, MAMMA1001633, MAMMA1001743, MAMMA1001837, MAMMA1002617, MAMMA1002869, MAMMA1002937, MAMMA1003011, NT2RM1000086, NT2RM1000187, NT2RM1000666, NT2RM1000885, NT2RM1000894, NT2RM1001059, NT2RM1001092, NT2RM2000013, NT2RM2000588, NT2RM2000624, NT2RM2000735, NT2RM2000740, NT2RM2001105, NT2RM2001635, NT2RM2001670, NT2RM2001771, NT2RM2001823, NT2RM2001936, NT2RM2001989, NT2RM2002004, NT2RM2002088, NT2RM2002091, NT2RM4000024, NT2RM4000046, NT2RM4000104, NT2RM4000202, NT2RM4000215, NT2RM4000290, NT2RM4000531, NT2RM4000751, NT2RM4000996, NT2RM4001092, NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001566, NT2RM4001592, 【1082】NT2RM4001597, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RP1000035, NT2RP1000111, NT2RP1000493, NT2RP1000574, NT2RP1000630, NT2RP1000902, NT2RP1000915, NT2RP1000958, NT2RP1000966, NT2RP1001013, NT2RP1001177, NT2RP2000008, NT2RP2000076, NT2RP2000126, NT2RP20

00153, NT2RP2000161, NT2RP2000248, NT2RP2000258, NT2RP2000297, NT2RP2000420, NT2RP2000931, NT2RP2001233, NT2RP2001420, NT2RP2001756, NT2RP2001869, NT2RP2002079, NT2RP2002270, NT2RP2002503, NT2RP2002591, NT2RP2002880, NT2RP2002939, NT2RP2002993, NT2RP2003137, NT2RP2003157, NT2RP2003277, NT2RP2003286, NT2RP2003308, NT2RP2003347, NT2RP2003714, NT2RP2003912, NT2RP2004013, NT2RP2004187, NT2RP2004689, NT2RP2004920, NT2RP2005393, NT2RP2005436, NT2RP2005496, NT2RP2005539, NT2RP2005701, NT2RP2005767, NT2RP2005776, NT2RP2005933, NT2RP2005942, NT2RP2006043, NT2RP2006436, NT2RP3000031, NT2RP3000050, NT2RP3000397, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP3000603, NT2RP3000632, NT2RP3000917, NT2RP3001057, NT2RP3001107, NT2RP3001120, NT2RP3001253, NT2RP3001338, NT2RP3001384, NT2RP3001398, NT2RP3001427, NT2RP3001428, NT2RP3001472, NT2RP3001646, NT2RP3001671, NT2RP3001792, NT2RP3001855, NT2RP3002056, NT2RP3002165, NT2RP3002399, NT2RP3002876, NT2RP3003193, NT2RP3003212, NT2RP3003555, NT2RP3004016, NT2RP3004206, NT2RP3004424, NT2RP3004428, NT2RP3004566, NT2RP3004617, NT2RP4000078, NT2RP4000111, NT2RP4000210,

【1083】NT2RP4000398, NT2RP4000481, NT2RP4000518, NT2RP4000997, NT2RP4001148, NT2RP4001206, NT2RP4001213, NT2RP4001433, NT2RP4001568, NT2RP4001638, NT2RP4001696, NT2RP4001753, NT2RP4001938, NT2RP4002058, NT2RP4002078, NT2RP4002081, NT2RP4002791, OVARC1000006, OVARC1000087, OVARC1000091, OVARC1000241, OVARC1000326, OVARC1000556, OVARC1000846, OVARC1001038, OVARC1001180, OVARC1001232, OVARC1001271, OVARC1001306, OVARC1001436, OVARC1002112, PLACE1000133, PLACE1000184, PLACE1000406, PLACE1000583, PLACE1000596, PLACE1000979, PLACE1001118, PLACE1001383, PLACE1001632, PLACE1002171, PLACE1002433, PLACE1002438, PLACE1002532, PLACE1002775, PLACE1002816, PLACE1002834, PLACE1003100, PLACE1003190, PLACE1003302, PLACE1003519, PLACE1003521, PLACE1003605, PLACE1003704, PLACE1003738, PLACE1003885, PLACE1003923, PLACE1004302, PLACE1004471, PLACE1004564, PLACE1004814, PLACE1004902, PLACE1005287, PLACE1005876, PLACE1005966, PLACE1006167, PLACE1006438, PLACE1006482, PLACE1006829, PLACE1006878, PLACE1006917, PLACE1007014, PLACE1007547, PLACE1007598, PLACE1007688, PLACE1007969, PLACE1008044, PLACE1008132, PLACE1008603, PLACE1009099, PLACE1009130, PLACE1009308, PLACE1009398, PLACE1010134, PLACE1010194, PLACE1010702, PLACE1010720, PLACE1010870, PLACE1011056, PLACE1011433, PLACE1011664, PLACE2000014, PLACE2000427, PLACE3000009, PLACE3000169, PLACE40

00014, PLACE4000156, PLACE4000192, PLACE4000261, PLACE4000326,

【1084】PLACE4000489, SKNMC1000011, THYR01000085, THYR01000242, THYR01000585, THYR01001100, THYR01001189, THYR01001809, Y79AA1000037, Y79AA1000214, Y79AA1000231, Y79AA1000589, Y79AA1000752, Y79AA1001391, Y79AA1001613, Y79AA1001705, Y79AA1001963, Y79AA1002431, Y79AA1002472, Y79AA1002482

【1085】DNA・RNA結合タンパク質のカテゴリに属すると推定されたクローンは、以下の292クローンであった。

HEMBA1000158, HEMBA1000216, HEMBA1000561, HEMBA1000591, HEMBA1000851, HEMBA1001088, HEMBA1001137, HEMBA1001405, HEMBA1001510, HEMBA1001804, HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1001869, HEMBA1002177, HEMBA1002935, HEMBA1003408, HEMBA1003545, HEMBA1003568, HEMBA1003591, HEMBA1003662, HEMBA1003684, HEMBA1003760, HEMBA1003783, HEMBA1003805, HEMBA1003953, HEMBA1004321, HEMBA1004354, HEMBA1004389, HEMBA1004479, HEMBA1004669, HEMBA1004847, HEMBA1004973, HEMBA1005202, HEMBA1005359, HEMBA1005931, HEMBA1006248, HEMBA1006278, HEMBA1006283, HEMBA1006359, HEMBA1006652, HEMBA1007087, HEMBA1007194, HEMBB1000264, HEMBB1000789, HEMBB1001011, HEMBB1001482, HEMBB1001736, HEMBB1001749, HEMBB1001839, HEMBB1002217, MAMMA1000183, MAMMA1000284, MAMMA1000731, MAMMA1001105, MAMMA1001222, MAMMA1001260, MAMMA1001743, MAMMA1001837, MAMMA1002385, MAMMA1002617, MAMMA1002869, MAMMA1002937, MAMMA1003011, NT2RM1000086, NT2RM1000539, NT2RM1000555, NT2RM1000666, NT2RM1000691, NT2RM1000826, NT2RM1000885, NT2RM1001059, NT2RM1001092, NT2RM2000371, NT2RM2000624, NT2RM2000735, NT2RM2001105, NT2RM2001424, NT2RM2001575, NT2RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001823, NT2RM2001989, NT2RM2002004, NT2RM2002014, NT2RM2002088, NT2RM2002091, NT2RM4000046, NT2RM4000104, NT2RM4000167, NT2RM4000191, NT2RM4000202, NT2RM4000531, NT2RM4000595, NT2RM4000733, NT2RM4000751, NT2RM4000996, NT2RM4001092, NT2RM4001140,

【1086】NT2RM4001178, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001880, NT2RM4001979, NT2RM4002093, NT2RM4002109, NT2RP1000470, NT2RP1000493, NT2RP1000574, NT2RP1000902, NT2RP1000966, NT2RP1001013, NT2RP1001073, NT2RP1001080, NT2RP2000008, NT2RP2000153, NT2RP2000258, NT2RP2000297, NT2RP2001127, NT2RP2001174, NT2RP2001233, NT2RP2001511, NT2RP2001756, NT2RP2001869, NT2RP2002079, NT2RP2002099, NT2RP2002503, NT2RP2002591, NT2RP2002939, NT2RP2003157, NT2RP2003329, NT2RP2003347, NT2RP2

2831

003480, NT2RP2003522, NT2RP2003564, NT2RP2003714, NT2RP2004187, NT2RP2004568, NT2RP2004920, NT2RP2005003, NT2RP2005139, NT2RP2005168, NT2RP2005436, NT2RP2005496, NT2RP2005701, NT2RP2005763, NT2RP2005776, NT2RP2005942, NT2RP2006043, NT2RP2006436, NT2RP2006464, NT2RP3000050, NT2RP30000512, NT2RP30000527, NT2RP30000562, NT2RP30000590, NT2RP30000603, NT2RP30000624, NT2RP30000632, NT2RP30000994, NT2RP30001057, NT2RP30001107, NT2RP30001120, NT2RP30001150, NT2RP30001155, NT2RP30001338, NT2RP30001398, NT2RP30001472, NT2RP30001672, NT2RP30001688, NT2RP30001724, NT2RP30001792, NT2RP30001855, NT2RP30002165, NT2RP30002399, NT2RP30002876, NT2RP30003138, NT2RP30003193, NT2RP30003251, NT2RP30003327, NT2RP30003555, NT2RP30004013, NT2RP30004078, NT2RP30004428, NT2RP30004490, NT2RP30004566, NT2RP30004594, NT2RP30004617, NT2RP30004618, NT2RP40000111, NT2RP40000398, NT2RP40000455, NT2RP40000518, NT2RP40000648,

【1087】NT2RP4000865, NT2RP4000929, NT2RP4001080, NT2RP4001095, NT2RP4001213, NT2RP4001433, NT2RP4001568, NT2RP4001696, NT2RP4001753, NT2RP4001838, NT2RP4001938, NT2RP4002078, OVARC1000006, OVARC10000087, OVARC1000241, OVARC1000746, OVARC10000846, OVARC1001232, OVARC1001271, OVARC1001306, OVARC1001987, OVARC1002112, PLACE1000406, PLACE1000583, PLACE1000979, PLACE1001118, PLACE1001632, PLACE1001739, PLACE1002438, PLACE1002532, PLACE1002775, PLACE1002834, PLACE1003302, PLACE1003519, PLACE1003605, PLACE1003704, PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004564, PLACE1004814, PLACE1005584, PLACE1005876, PLACE1005951, PLACE1006196, PLACE1006482, PLACE1006488, PLACE1006531, PLACE1006917, PLACE1007346, PLACE1007547, PLACE1007598, PLACE1007688, PLACE1007969, PLACE1008132, PLACE1009099, PLACE1009246, PLACE1009398, PLACE1009476, PLACE1009622, PLACE1010053, PLACE1010194, PLACE1010702, PLACE1010870, PLACE1011056, PLACE1011114, PLACE1011433, PLACE2000427, PLACE3000009, PLACE3000169, PLACE4000014, PLACE4000156, PLACE4000192, PLACE4000261, PLACE4000489, SKNMC1000091, THYR01000085, THYR01000242, THYR01000501, THYR01001100, THYR01001189, THYR01001809, Y79AA1000037, Y79AA1000349, Y79AA1000752, Y79AA1001211, Y79AA1001312, Y79AA1001391, Y79AA1001613, Y79AA1002103, Y79AA1002472, Y79AA1002482, RNA合成関連タンパク質のカテゴリに属すると推定されたクローンは、以下の66クローンであった。

HEMBA1000591, HEMBA1001579, HEMBA1003179, HEMBA1003591, HEMBA1006278, HEMBB1000226, NT2RM1000187, NT2RM1000852, NT2RM2000624, NT2RM2001989, NT2RM2002100, NT2RM4000191, NT2RM4001178, NT2RM4002093, NT2R

2832

P1000035, NT2RP1000272, NT2RP1000470, NT2RP1001080, NT2RP2000153, NT2RP2002928, NT2RP2003157, NT2RP2004568, NT2RP2005126, NT2RP2005436, NT2RP2005539, NT2RP2005605, NT2RP2005776, NT2RP2005942, NT2RP2006043, NT2RP2006238, NT2RP3000361, NT2RP3000397, NT2RP3001671, NT2RP3004504, NT2RP4000078, NT2RP4000111, NT2RP4000481, NT2RP4000518, NT2RP4000614, NT2RP4000929, NT2RP4001696, NT2RP4002058, OVARC1001232, OVARC1001577, PLACE1000406, PLACE1000596, PLACE1000755, PLACE1001739, PLACE1003704, PLACE1003885, PLACE1004564, PLACE1004814, PLACE1004902, PLACE1005373, PLACE1005646, PLACE1005876, PLACE1006196, PLACE1006626, PLACE1006878, PLACE1006917, PLACE1009476, PLACE1009925, PLACE1010194, PLACE1011114, THYR01000121, Y79AA1001963,

【1088】タンパク質合成・輸送関連タンパク質のカテゴリに属すると推定されたクローンは以下の183クローンであった。

HEMBA1000012, HEMBA1000141, HEMBA1000592, HEMBA1003617, HEMBA1003773, HEMBA1004202, HEMBA1004276, HEMBA1004734, HEMBA1004847, HEMBA1004929, HEMBA1004930, HEMBA1005047, HEMBA1005202, HEMBA1006031, HEMBA1006272, HEMBA1006474, HEMBA1006652, HEMBA1006914, HEMBA1006973, HEMBA1007224, HEMBB1000915, HEMBB1001112, HEMBB1001137, HEMBB1001736, HEMBB1001831, HEMBB1001915, MAMMA1000085, MAMMA1000734, MAMMA1001008, MAMMA1002170, MAMMA1002219, MAMMA1002236, MAMMA1002619, NT2RM1000661, NT2RM1000833, NT2RM2000092, NT2RM2000504, NT2RM2000577, NT2RM2000821, NT2RM2001201, NT2RM2001592, NT2RM2001613, NT2RM2001648, NT2RM2001730, NT2RM2001760, NT2RM2002055, NT2RM4000155, NT2RM4000169, NT2RM4000344, NT2RM4000356, NT2RM4000421, NT2RM4000712, NT2RM4001054, NT2RM4001203, NT2RM4001382, NT2RM4001444, NT2RM4002062, NT2RM4002205, NT2RM4002623, NT2RP1000326, NT2RP1000522, NT2RP1000547, NT2RP1000746, NT2RP1000947, NT2RP1001569, NT2RP2000147, NT2RP2000710, NT2RP2000880, NT2RP2000943, NT2RP2001290, NT2RP2001392, NT2RP2001601, NT2RP2001613, NT2RP2001660, NT2RP2001740, NT2RP2002124, NT2RP2002606, NT2RP2002862, NT2RP2002959, NT2RP2002980, NT2RP2003137, NT2RP2003158, NT2RP2003391, NT2RP2003394, NT2RP2003401, NT2RP2003433, NT2RP2003704, NT2RP2003713, NT2RP2003737, NT2RP2003760, NT2RP2003981, NT2RP2004366, NT2RP2004389, NT2RP2004791, NT2RP2005012, NT2RP2005116, NT2RP2005360, NT2RP2005763, NT2RP2005784, NT2RP3000366,

【1089】NT2RP3000759, NT2RP3000968, NT2RP3001113, NT2RP3001690, NT2RP3002045, NT2RP3002151, NT2RP3002529, NT2RP3002671, NT2RP3003301, NT2RP3003846, NT2RP3003876, NT2RP3004209, NT2RP4000370, NT2RP

2833

4000457, NT2RP4000879, NT2RP4000927, NT2RP4001041, NT2RP4001117, NT2RP4001313, NT2RP4001315, NT2RP4001574, NT2RP4001592, OVARC1000013, OVARC1000071, OVARC1000085, OVARC1000465, OVARC1000564, OVARC1000771, OVARC1000862, OVARC1001171, OVARC1001180, OVARC1001342, PLACE1000007, PLACE1000061, PLACE1000081, PLACE1000492, PLACE1000863, PLACE1001092, PLACE1001748, PLACE1002090, PLACE1003174, PLACE1003915, PLACE1004104, PLACE1004270, PLACE1004743, PLACE1005557, PLACE1005813, PLACE1006170, PLACE1006488, PLACE1006829, PLACE1007706, PLACE1007729, PLACE1008273, PLACE1008402, PLACE1008790, PLACE1008813, PLACE1009094, PLACE1009130, PLACE1009477, PLACE1009721, PLACE1009845, PLACE1010074, PLACE1010547, PLACE1011109, PLACE1011229, PLACE1011477, PLACE1012031, PLACE2000404, PLACE3000059, PLACE3000121, PLACE4000269, PLACE4000654, SKNMC1000011, THYRO1000983, THYRO1001003, THYRO1001313, Y79AA1000560, Y79AA1000784, Y79AA1000968, Y79AA1001493, Y79AA1001875, Y79AA1002027, Y79AA1002209,

【1090】細胞骨格関連タンパク質のカテゴリーに属すると推定されたクローンは、以下の130クローンであった。

HEMBA1000156, HEMBA1000168, HEMBA1000411, HEMBA1000588, HEMBA1001043, HEMBA1001651, HEMBA1001661, HEMBA1002102, HEMBA1002161, HEMBA1002939, HEMBA1003235, HEMBA1003581, HEMBA1004499, HEMBA1004534, HEMBA1004697, HEMBA1004929, HEMBA1004972, HEMBA1005582, HEMBA1005595, HEMBA1006344, HEMBA1006737, HEMBB1001175, HEMBB1001282, HEMBB1001562, HEMBB1001802, MAMMA1000824, MAMMA1001041, MAMMA1001576, MAMMA1001679, MAMMA1001735, MAMMA1002297, MAMMA1002351, MAMMA1002622, MAMMA1002637, MAMMA1003127, NT2RM1000850, NT2RM1000898, NT2RM2000030, NT2RM2000260, NT2RM2000691, NT2RM2001324, NT2RM4000169, NT2RM4000229, NT2RM4000515, NT2RM4001217, NT2RP1000202, NT2RP1000348, NT2RP1000460, NT2RP1000478, NT2RP1001033, NT2RP1001294, NT2RP1001302, NT2RP2000070, NT2RP2000812, NT2RP2000814, NT2RP2001168, NT2RP2001245, NT2RP2001634, NT2RP2001900, NT2RP2003307, NT2RP2003394, NT2RP2004041, NT2RP2004242, NT2RP2004538, NT2RP2004587, NT2RP2004681, NT2RP2004732, NT2RP2004978, NT2RP2005491, NT2RP2005531, NT2RP2005712, NT2RP2006275, NT2RP3000753, NT2RP3001113, NT2RP3001216, NT2RP3001239, NT2RP3001272, NT2RP3001554, NT2RP3001690, NT2RP3001799, NT2RP3002688, NT2RP3003061, NT2RP3003185, NT2RP3003230, NT2RP3004569, NT2RP3004578, NT2RP4001004, NT2RP4001086, NT2RP4001256, NT2RP4001567, NT2RP4001927, OVARC1000001, OVARC1000106, OVARC1000437, OVARC1000520, OVARC1000679, OVARC

2834

1001731, OVARC1002050, PLACE1001104, PLACE1002571, PLACE1002591, PLACE1002655, PLACE1002714, PLACE1003625, PLACE1005287, PLACE1006552, PLACE1007946, PLACE1008426, PLACE1010148, PLACE1010547, PLACE1010743, PLACE1010896, PLACE1010960, PLACE1011310, PLACE1011922, PLACE2000216, PLACE2000274, PLACE2000371, PLACE2000458, PLACE3000145, PLACE3000416, PLACE4000009, THYRO1000132, THYRO1001405, THYRO1001458, Y79AA1000368, Y79AA1000794, Y79AA1000833, Y79AA1000962, Y79AA1002208,

【1091】細胞分裂・増殖関連タンパク質のカテゴリーに属すると推定されたクローンは、以下の54クローンであった。

HEMBA1001019, HEMBA1001595, HEMBA1002363, HEMBA1002997, HEMBA1003136, HEMBA1003369, HEMBA1004131, HEMBA1004354, HEMBA1005621, HEMBB1000037, HEMBB1000264, MAMMA1001768, MAMMA1002769, NT2RM1000354, NT2RM1000430, NT2RM1000874, NT2RM2001256, NT2RM2001743, NT2RM2001896, NT2RM2002145, NT2RM4000215, NT2RM4001714, NT2RP1000163, NT2RP1000333, NT2RP1000439, NT2RP2000346, NT2RP2001397, NT2RP2002595, NT2RP2003177, NT2RP2003596, NT2RP2003912, NT2RP2004396, NT2RP2005037, NT2RP2005520, NT2RP2005669, NT2RP2005835, NT2RP3001730, NT2RP3002081, NT2RP4000210, NT2RP4000415, NT2RP4001414, NT2RP4001634, OVARC10000013, OVARC1000937, PLACE1001383, PLACE1002433, PLACE1004316, PLACE1005287, PLACE1008808, PLACE1010720, PLACE1010833, Y79AA1000748, Y79AA1001236, Y79AA1001394,

【1092】発生・発達関連タンパク質のカテゴリーに属すると推定されたクローンは、以下の36クローンであった。

HEMBA1000518, HEMBA1001847, HEMBA1001869, HEMBA1003545, HEMBA1004973, HEMBB1002442, MAMMA1001837, NT2RM2001670, NT2RM4000046, NT2RM4000531, NT2RM4001140, NT2RM4001858, NT2RP2002078, NT2RP2004187, NT2RP2006436, NT2RP3000603, NT2RP3000994, NT2RP3001580, NT2RP3001708, NT2RP3003071, NT2RP3004472, NT2RP3004617, NT2RP4000246, NT2RP4001567, OVARC1000304, OVARC1000746, PLACE1000793, PLACE1002532, PLACE1003258, PLACE1003625, PLACE1004460, PLACE1009622, PLACE4000558, THYRO1000085, Y79AA1001391, Y79AA1001692,

【1093】細胞防御関連タンパク質のカテゴリーに属すると推定されたクローンは以下の30クローンであった。

HEMBA1000005, HEMBA1000531, HEMBA1003417, HEMBA1006253, NT2RM4000354, NT2RM4001880, NT2RP1000333, NT2RP1000493, NT2RP2000006, NT2RP2000045, NT2RP2000809, NT2RP2001536, NT2RP2002464, NT2RP2004920, NT2R

2835

P2005037, NT2RP3000590, NT2RP3001426, NT2RP3002062, NT2RP3002785, NT2RP3004262, NT2RP4001555, NT2RP4001638, PLACE1006958, PLACE1008275, PLACE1009113, PLACE1011858, PLACE4000014, THYR01000684, Y79AA1002139, Y79AA1002229,

【1094】上記以外のクローンのうち以下の261クローンについては、上記のいずれのカテゴリに属するか明らかでないクローンであったが、全長配列に対する相同性検索で何らかの機能が予測されているクローンである。クローン名と相同性検索結果のDefinitionを//で区切り、以下に示した。

HEMBA1000030//Homo sapiens ARF GTPase-activating protein GIT1 mRNA, complete cds.

HEMBA1000307//CARNITINE DEFICIENCY-ASSOCIATED PROTEIN EXPRESSED IN VENTRICLE 1

HEMBA1000333//Homo sapiens F-box protein Fbx21 (FBX21) mRNA, complete cds.

HEMBA1000488//RING CANAL PROTEIN (KELCH PROTEIN).

HEMBA1000523//TESTIS-SPECIFIC PROTEIN PBS13.

HEMBA1001197//Homo sapiens rap2 interacting protein mRNA, complete cds.

HEMBA1001302//Homo sapiens calcium binding protein precursor, mRNA, complete cds.

HEMBA1001455//Mus musculus transposon-derived Buster2 transposase-like protein gene, partial cds.

HEMBA1001675//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS9.

HEMBA1001714//Homo sapiens mRNA for ATPase inhibitor or precursor, complete cds.

HEMBA1001744//SCY1 PROTEIN.

HEMBA1001967//Homo sapiens NY-REN-57 antigen mRNA, partial cds.

HEMBA1002151//Rattus norvegicus p34 mRNA, complete cds.

HEMBA1002215//TESTIN 2 (TES2) [CONTAINS: TESTIN 1 (TES1)].

HEMBA1002458//OVARIAN GRANULOSA CELL 13.0 KD PROTEIN HGR74.

HEMBA1002777//Fugu rubripes BAW (BAW) mRNA, complete cds.

HEMBA1003098//Homo sapiens NY-REN-6 antigen mRNA, partial cds.

HEMBA1003199//Homo sapiens chromosome 5 F-box protein Fbx4 (FBX4) mRNA, complete cds.

HEMBA1003615//Homo sapiens ART-4 mRNA, complete cds.

HEMBA1003836//MOB1 PROTEIN (MPS1 BINDER 1).

HEMBA1004295//Homo sapiens NY-REN-25 antigen mRNA, partial cds.

HEMBA1004573//Homo sapiens mRNA for HELG protein.

2836

HEMBA1004604//Homo sapiens COP9 complex subunit 7a mRNA, complete cds.

HEMBA1004795//CDC4-LIKE PROTEIN (FRAGMENT).

HEMBA1005101//Homo sapiens SYT interacting protein SIP mRNA, complete cds.

HEMBA1005201//Homo sapiens CGI-07 protein mRNA, complete cds.

HEMBA1005206//Drosophila simulans anon73B1 gene and Su(P) gene.

10 HEMBA1005530//Homo sapiens anaphase-promoting complex subunit 7 (APC7) mRNA, complete cds.

HEMBA1005666//Homo sapiens mRNA for DIPB protein.

HEMBA1005990//Homo sapiens I-1 receptor candidate protein mRNA, complete cds.

HEMBA1006268//Homo sapiens HQ0024c mRNA, complete cds.

HEMBA1006398//Human L1 element L1.6 putative p150 gene, complete cds.

HEMBA1006445//Homo sapiens putative tumor suppressor NOEY2 mRNA, complete cds.

HEMBA1007174//Homo sapiens epsin 2b mRNA, complete cds.

HEMBA1007251//Homo sapiens F-box protein FBX29 (FBX29) mRNA, partial cds.

HEMBA1000036//Homo sapiens CGI-51 protein mRNA, complete cds.

HEMBA1000144//GUANYLATE CYCLASE ACTIVATING PROTEIN 2 (GCAP 2) (RETINAL GUANYLYL CYCLASE ACTIVATOR PROTEIN P24).

30 HEMBA1000973//Mus musculus schlafen3 (Slfn3) mRNA, complete cds.

HEMBA1001058//Homo sapiens neuronal thread protein AD7c-NTP mRNA, complete cds

HEMBA1001234//65 KD YES-ASSOCIATED PROTEIN (YAP65).

HEMBA1001288//COPPER HOMEOSTASIS PROTEIN CUTC.

HEMBA1001331//Mus musculus mRNA for hepatoma-derived growth factor, complete cds, strain: BALB/c.

40 HEMBA1001384//Homo sapiens COP9 complex subunit 4 mRNA, complete cds.

HEMBA1002266//NEURONAL PROTEIN.

HEMBA1002510//GYP7 PROTEIN.

HEMBA1002705//Homo sapiens CGI-27 protein mRNA, complete cds.

MAMMA1000055//TESTIN 2 (TES2) [CONTAINS: TESTIN 1 (TES1)].

MAMMA1000625//GYP7 PROTEIN.

MAMMA1001075//Homo sapiens CGI-72 protein mRNA, complete cds.

50 MAMMA1001181//ABC1 PROTEIN HOMOLOG PRECURSOR.

2837

MAMMA1001259//Mus musculus F-box protein FBX18 mRNA, partial cds.
 MAMMA1001730//Homo sapiens brain and nasopharyngeal carcinoma susceptibility protein NSG-x mRNA, partial cds.
 MAMMA1002143//Homo sapiens Cdc42 effector protein 4 mRNA, complete cds.
 MAMMA1002699//Rattus norvegicus EH domain binding protein Epsin mRNA, complete cds.
 MAMMA1002972//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS27.
 MAMMA1003113//Mus musculus COP9 complex subunit 7a (COPS7a) mRNA, complete cds.
 NT2RM1000118//CALCINEURIN B SUBUNIT (PROTEIN PHOSPHATASE 2B REGULATORY SUBUNIT) (CALCINEURIN REGULATORY SUBUNIT).
 NT2RM1000186//CALCINEURIN B SUBUNIT (PROTEIN PHOSPHATASE 2B REGULATORY SUBUNIT) (CALCINEURIN REGULATORY SUBUNIT).
 NT2RM1000244//Homo sapiens TRAF4 associated factor 1 mRNA, partial cds.
 NT2RM1000421//RIBONUCLEASE INHIBITOR.
 NT2RM1000499//Caenorhabditis elegans mRNA for centaurin gamma 1A.
 NT2RM1000623//RIBONUCLEASE INHIBITOR.
 NT2RM1000883//Homo sapiens I-1 receptor candidate protein mRNA, complete cds.
 NT2RM2000502//Rattus norvegicus W307 mRNA, complete cds.
 NT2RM2000599//Homo sapiens F-box protein Lilina (LILINA) mRNA, complete cds.
 NT2RM2000718//Homo sapiens endocrine regulator mRNA, complete cds.
 NT2RM2001065//Homo sapiens COP9 complex subunit 4 mRNA, complete cds.
 NT2RM2001196//PROLINE-RICH PROTEIN MP-3 (FRAGMENT).
 NT2RM2001983//Homo sapiens RGS-GAIP interacting protein GIPC mRNA, complete cds.
 NT2RM2002109//Homo sapiens glioma amplified on chromosome 1 protein (GAC1) mRNA, complete cds.
 NT2RM2002142//GASTRULATION SPECIFIC PROTEIN G12.
 NT2RM4000030//LAS1 PROTEIN.
 NT2RM4000139//R. norvegicus trg mRNA.
 NT2RM4000156//H. sapiens HPBRII-7 gene.
 NT2RM4000386//Mus musculus ODZ3 (Odz3) mRNA, partial cds.
 NT2RM4000590//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RM4001047//MO25 PROTEIN.
 NT2RM4001155//ADRENAL MEDULLA 50 KD PROTEIN.

2838

NT2RM4001256//Xenopus laevis putative Zic3 binding protein mRNA, complete cds.
 NT2RM4001320//Homo sapiens mRNA for Neuroblastoma, complete cds.
 NT2RM4001340//UTR4 PROTEIN (UNKNOWN TRANSCRIPT 4 PROTEIN).
 NT2RM4001347//Homo sapiens NY-REN-25 antigen mRNA, partial cds.
 NT2RM4001371//Homo sapiens IDN3 mRNA, partial cds.
 NT2RM4001582//Mus musculus COP9 complex subunit 7b (COPS7b) mRNA, complete cds.
 NT2RM4001611//SIS2 PROTEIN (HALOTOLERANCE PROTEIN HAL3).
 NT2RM4001731//Homo sapiens F-box protein Lilina (LILINA) mRNA, complete cds.
 NT2RM4001969//R. norvegicus mRNA for IP63 protein.
 NT2RM4002034//Homo sapiens hiwi mRNA, partial cds.
 NT2RM4002075//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RM4002226//GTPASE ACTIVATING PROTEIN ROTUND.
 NT2RP1000040//Mus musculus donson protein (Donson) mRNA, partial cds.
 NT2RP1000363//R. norvegicus LL5 mRNA.
 NT2RP1000481//Homo sapiens antigen NY-CO-3 (NY-CO-3) mRNA, partial cds.
 NT2RP1000513//Human NifU-like protein (hNifU) mRNA, partial cds.
 NT2RP1000733//Human mRNA for GSPT1-TK protein, complete cds.
 NT2RP1000860//Homo sapiens KLO4P mRNA, complete cds.
 NT2RP1000954//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP1001011//Drosophila melanogaster putative 43 kDa protein (TH1) mRNA, complete cds.
 NT2RP1001395//Homo sapiens COP9 complex subunit 7a mRNA, complete cds.
 NT2RP1001457//Homo sapiens partial mRNA for beta-transducin family protein (putative).
 NT2RP1001494//MALE STERILITY PROTEIN 2.
 NT2RP2000054//Homo sapiens putative ring zinc finger protein NY-REN-43 antigen mRNA, complete cds.
 NT2RP2000067//Mus musculus ODZ3 (Odz3) mRNA, partial cds.
 NT2RP2000133//Homo sapiens Leman coiled-coil protein (LCCP) mRNA, complete cds.
 NT2RP2000157//MLO2 PROTEIN.
 NT2RP2000764//NIFS PROTEIN.
 NT2RP2000965//Homo sapiens mRNA for fls353, complete cds.
 NT2RP2001839//SCY1 PROTEIN.
 NT2RP2001883//Homo sapiens CGI-01 protein mRNA, co

- mplete cds.
 NT2RP2001976//Mus musculus calmodulin-binding protein SHA1 (Shal) mRNA, complete cds.
 NT2RP2001985//Homo sapiens high-risk human papilloma viruses E6 oncoproteins targeted protein E6TP1 alpha mRNA, complete cds.
 NT2RP2002185//Homo sapiens ubiquitin mRNA, complete cds.
 NT2RP2002442//HESA PROTEIN.
 NT2RP2002727//Rattus norvegicus tulip 2 mRNA, complete cds. 10
 NT2RP2002741//Homo sapiens mRNA for Neuroblastoma, complete cds.
 NT2RP2002986//Homo sapiens mRNA for Kelch motif containing protein, complete cds.
 NT2RP2003121//Mus musculus enhancer of polycomb (Epc1) mRNA, complete cds.
 NT2RP2003265//Homo sapiens CGI-53 protein mRNA, complete cds.
 NT2RP2003272//Homo sapiens ubiquitin mRNA, complete cds. 20
 NT2RP2003857//MYOTROPHIN (V-1 PROTEIN) (GRANULE CELL DIFFERENTIATION PROTEIN).
 NT2RP2003871//Homo sapiens transposon-derived Buster1 transposase-like protein gene, complete cds.
 NT2RP2004425//Mus musculus axotrophin mRNA, complete cds.
 NT2RP2004476//Homo sapiens cyclin L ania-6a mRNA, complete cds.
 NT2RP2004710//Mus musculus formin binding protein 30 mRNA, complete cds. NT2RP2004816//H58 PROTEIN.
 NT2RP2005441//Homo sapiens hypothalamus protein HT002 mRNA, complete cds.
 NT2RP2005490//Mus musculus D3M3e (D3M3e) mRNA, complete cds.
 NT2RP2005620//Homo sapiens epsin 2a mRNA, complete cds.
 NT2RP2005654//CYSTEINE STRING PROTEIN (CCCS1).
 NT2RP2005675//Homo sapiens growth suppressor related (DOC-1R) mRNA, complete cds.
 NT2RP2005753//Homo sapiens I-1 receptor candidate protein mRNA, complete cds. 40
 NT2RP2005841//Homo sapiens mRNA for ALEX3, complete cds.
 NT2RP2006598//Homo sapiens retinoid x receptor interacting protein mRNA, complete cds.
 NT2RP3000047//NPL4 PROTEIN.
 NT2RP3000233//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP3000868//Human ovarian cancer downregulated myosin heavy chain homolog (Doc1) mRNA, complete cds. 50
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 NT2RP3000869//Drosophila melanogaster AAA family protein Bor (bor) mRNA, complete cds.
 NT2RP3001399//SSU72 PROTEIN.
 NT2RP3001407//SCY1 PROTEIN.
 NT2RP3001457//Drosophila melanogaster Melted (melt) mRNA, partial cds.
 NT2RP3001587//Human anthracycline-associated resistance ARX mRNA, complete cds.
 NT2RP3001712//Homo sapiens HP1-BP74 protein mRNA, complete cds.
 NT2RP3001819//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP3001854//Homo sapiens novel retinal pigment epithelial cell protein (NORPEG) mRNA, complete cds.
 NT2RP3001931//Rattus norvegicus clone C48 CDK5 activator-binding protein mRNA, complete cds.
 NT2RP3002273//SCD6 PROTEIN.
 NT2RP3002631//Homo sapiens Ran binding protein 11 mRNA, complete cds.
 NT2RP3002682//Homo sapiens CGI-145 protein mRNA, complete cds.
 NT2RP3002770//MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD116.
 NT2RP3002818//INSERTION ELEMENT IS2A HYPOTHETICAL 48.2 KD PROTEIN.
 NT2RP3002948//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP3002972//Halocynthia roretzi mRNA for HrPET-1, complete cds.
 NT2RP3003032//Homo sapiens okadaic acid-inducible and cAMP-regulated phosphoprotein 19 (ARPP-19) mRNA, complete cds.
 NT2RP3003290//Mus musculus mRNA for Ndr1 related protein Ndr3, complete cds.
 NT2RP3003411//Mus musculus COP9 complex subunit 7b (COPS7b) mRNA, complete cds.
 NT2RP3003491//Drosophila melanogaster Pelle associated protein Pellino (Pli) mRNA, complete cds.
 NT2RP3003500//SCY1 PROTEIN.
 NT2RP3003726//Homo sapiens spermatogenesis associated PD1 mRNA, complete cds.
 NT2RP3004348//R. norvegicus mRNA for cytosolic resistiniferatoxin-binding protein.
 NT2RP3004507//MOB1 PROTEIN (MPS1 BINDER 1).
 NT2RP4000129//Xenopus laevis F-box protein 28 (Fbx28) mRNA, partial cds. NT2RP4000498//MOB1 PROTEIN (MPS1 BINDER 1).
 NT2RP4000528//NPL4 PROTEIN.
 NT2RP4000737//Mus musculus F-box protein FBL10 mRNA, partial cds.
 NT2RP4000979//Homo sapiens putative HIV-1 infectio

2841

n related protein mRNA, partial cds.
 NT2RP4001010//Rattus norvegicus PSD-95/SAP90-associated protein-4 mRNA, complete cds.
 NT2RP4001207//Homo sapiens Ran binding protein 11 mRNA, complete cds.
 NT2RP4001228//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP4001260//Homo sapiens F-box protein Fbx21 (FBX21) mRNA, complete cds.
 NT2RP4001339//Homo sapiens mRNA for AMMERC1 protein.
 NT2RP4001351//Human ovarian cancer downregulated myosin heavy chain homolog (Doc1) mRNA, complete cds.
 NT2RP4001474//Xenopus laevis putative Zic3 binding protein mRNA, complete cds.
 NT2RP4001966//Mus musculus ODZ3 (Odz3) mRNA, partial cds.
 NT2RP4002018//RING CANAL PROTEIN (KELCH PROTEIN).
 OVARC1000209//Oryza sativa submergence induced protein 2A mRNA, complete cds.
 OVARC1000876//MOB1 PROTEIN (MPS1 BINDER 1).
 OVARC1001065//Homo sapiens CGI-12 protein mRNA, complete cds.
 OVARC1001092//Homo sapiens mRNA for JM5 protein, complete CDS (clone IMAGE 53337, LLNLc110F1857Q7 (RZPD Berlin) and LLNLc110G0913Q7 (RZPD Berlin)).
 OVARC1001419//Homo sapiens GOK (STIM1) mRNA, complete cds.
 OVARC1001555//NGG1-INTERACTING FACTOR 3.
 OVARC1001711//CORNIFIN B (SMALL PROLINE-RICH PROTEIN 1B) (SPR1B) (SPR1 B).
 OVARC1001943//Mus musculus DEBT-91 mRNA, complete cds.
 PLACE1000004//Homo sapiens IDN3-B mRNA, complete cds.
 PLACE1000066//SSU72 PROTEIN.
 PLACE1000610//MSN5 PROTEIN.
 PLACE1000636//MALE STERILITY PROTEIN 2.
 PLACE1000769//Homo sapiens CGI-18 protein mRNA, complete cds.
 PLACE1000987//Rattus norvegicus late gestation lung 2 protein (Lgl2) mRNA, complete cds.
 PLACE1001036//Homo sapiens mRNA for alpha integrin binding protein 63, partial.
 PLACE1001845//Mus musculus cyclin ania-6a mRNA, complete cds.
 PLACE1001920//Homo sapiens MDC-3.13 isoform 2 mRNA, complete cds.
 PLACE1002665//Mus musculus enhancer of polycomb (Epc1) mRNA, complete cds.

2842

PLACE1003602//Homo sapiens mRNA expressed in placenta.
 PLACE1003611//Homo sapiens anaphase-promoting complex subunit 4 (APC4) mRNA, complete cds.
 PLACE1004256//Mus musculus short coiled coil protein in SCOC (Scoc) mRNA, complete cds.
 PLACE1004550//Homo sapiens CGI-20 protein mRNA, complete cds.
 PLACE1004868//MALE STERILITY PROTEIN 2.
 10 PLACE1004930//Homo sapiens MDC-3.13 isoform 2 mRNA, complete cds.
 PLACE1005052//Homo sapiens CGI-16 protein mRNA, complete cds.
 PLACE1005102//RING CANAL PROTEIN (KELCH PROTEIN).
 PLACE1005176//Homo sapiens hypothalamus protein HT001 mRNA, complete cds.
 PLACE1005187//APAG PROTEIN.
 PLACE1005331//Homo sapiens 7h3 protein mRNA, partial cds.
 20 PLACE1005727//Homo sapiens STRIN protein (STRIN) mRNA, complete cds.
 PLACE1006003//Homo sapiens CGI-94 protein mRNA, complete cds.
 PLACE1006335//Homo sapiens NY-REN-50 antigen mRNA, partial cds.
 PLACE1006385//Homo sapiens epsin 2a mRNA, complete cds.
 PLACE1006506//Homo sapiens anaphase-promoting complex subunit 4 (APC4) mRNA, complete cds.
 30 PLACE1007105//Homo sapiens muskelin (MKLN1) mRNA, complete cds.
 PLACE1007537//Homo sapiens ankyrin repeat-containing protein ASB-2 mRNA, complete cds.
 PLACE1007705//Mus musculus mRNA for Ndr1 related protein Ndr3, complete cds.
 PLACE1007791//Homo sapiens IDN3-B mRNA, complete cds.
 PLACE1007897//Homo sapiens FLASH mRNA, complete cds.
 40 PLACE1008080//Homo sapiens mRNA for HEXIM1 protein, complete cds.
 PLACE1008368//RING CANAL PROTEIN (KELCH PROTEIN).
 PLACE1008398//GENE 33 POLYPEPTIDE.
 PLACE1008465//Homo sapiens mRNA for rapa-1 (rapa gene).
 PLACE1008627//Homo sapiens mRNA for cysteine-rich protein.
 PLACE1009020//NIFS PROTEIN.
 PLACE1009060//BR01 PROTEIN.
 50 PLACE1009186//Homo sapiens small zinc finger-like

2843

protein (TIM9b) mRNA, complete cds.
 PLACE1009443//Mus musculus F-box protein FBL8 mRNA, complete cds.
 PLACE1009571//Homo sapiens PTD002 mRNA, complete cds.
 PLACE1009670//Homo sapiens genethonin 1 mRNA, complete cds.
 PLACE1010105//RING CANAL PROTEIN (KELCH PROTEIN).
 PLACE1010261//SEGREGATION DISTORTER PROTEIN.
 PLACE1010310//SPIDROIN 2 (DRAGLINE SILK FIBROIN 2) (FRAGMENT).
 PLACE1010522//Homo sapiens mRNA for DEPP (decidual protein induced by progesterone), complete cds.
 PLACE1010579//Homo sapiens CED-6 protein (CED-6) mRNA, complete cds.
 PLACE1010628//Homo sapiens S164 gene, partial cds; PS1 and hypothetical protein genes, complete cds; and S171 gene, partial cds.
 PLACE1010661//TESTIS-SPECIFIC PROTEIN PBS13.
 PLACE1010761//Homo sapiens mRNA for cisplatin resistance-associated overexpressed protein, complete cds.
 PLACE1011185//INSERTION ELEMENT IS1 PROTEIN INSB.
 PLACE1011340//Homo sapiens IDN3-B mRNA, complete cds.
 PLACE1011586//Rattus norvegicus clone C53 CDK5 activator-binding protein mRNA, complete cds.
 PLACE2000246//RING CANAL PROTEIN (KELCH PROTEIN).
 PLACE2000411//Homo sapiens epsin 2b mRNA, complete cds.
 PLACE3000477//Homo sapiens phosphoprotein pp75 mRNA, partial cds.
 THYR01000173//Homo sapiens AP-mu chain family member mu1B (HSMU1B) mRNA, complete cds.
 THYR01000401//Human Tcd37 homolog (HTcd37) mRNA, partial cds.
 THYR01000666//Mus musculus mRNA for kinesin like protein 9.
 THYR01001033//TRANSFORMATION-SENSITIVE PROTEIN IEF SSP 3521.
 THYR01001347//Homo sapiens RAN binding protein 16 mRNA, complete cds.
 THYR01001656//Homo sapiens Leman coiled-coil protein (LCCP) mRNA, complete cds.
 THYR01001703//NIFR3-LIKE PROTEIN.
 THYR01001721//RING CANAL PROTEIN (KELCH PROTEIN).
 Y79AA1000059//Homo sapiens aryl-hydrocarbon interacting protein-like 1 (AIP1) gene, complete cds.
 Y79AA1000181//Homo sapiens CGI-01 protein mRNA, complete cds.

2844

Y79AA1000268//Mus musculus Nip2l mRNA, complete cds.
 Y79AA1000313//CALPHOTIN.
 Y79AA1000540//CELL POLARITY PROTEIN TEA1.
 Y79AA1000966//Homo sapiens COP9 complex subunit 4 mRNA, complete cds.
 Y79AA1000985//Human centrosomal protein kendrin mRNA, complete cds.
 Y79AA1001323//Mus musculus mRNA for GSG1, complete cds.
 Y79AA1001402//Homo sapiens paraneoplastic cancer-testis-brain antigen (MA4) mRNA, partial cds.
 Y79AA1001679//Homo sapiens lambda-crystallin mRNA, complete cds.
 Y79AA1001923//Homo sapiens F-box protein Fbx22 (FBX22) gene, partial cds.
 Y79AA1002083//H. sapiens mRNA for MUF1 protein.
 Y79AA1002307//Homo sapiens astrotactin2 (ASTN2) mRNA, complete cds.
 Y79AA1002311//R. norvegicus mRNA for cytosolic resiniferatoxin-binding protein.
 Y79AA1002487//Homo sapiens chromosome 5 F-box protein Fbx4 (FBX4) mRNA, complete cds.
 【1095】前出以外のクローンのうち、Pfamによるドメイン検索の結果から機能予測され、上記と同様な機能カテゴリー分類が可能なクローンが36クローンあった。分類は以下の通りであった。分泌・膜タンパク質に属する可能性が高いクローンはMAMMA1002498, NT2RM4002287の2クローンであった。糖タンパク質関連タンパク質に属する可能性が高いクローンはMAMMA1002498の1クローンであった。シグナル伝達関連タンパク質に属する可能性が高いクローンはHEMBA1001247, NT2RM2001813, NT2RM4001454, NT2RP2005140, NT2RP2005293, NT2RP3000487, NT2RP3003311, PLACE1000972, PLACE1003723, PLACE1005327, PLACE3000124の11クローンであった。転写関連タンパク質に属する可能性が高いクローンはHEMBA1003257, NT2RM2000101, NT2RM2001797, NT2RP1000101, NT2RP2002208, NT2RP3001214, NT2RP3003278, NT2RP4001235, PLACE1000050, PLACE1001716, PLACE1002499, PLACE1007544の12クローンであった。酵素・代謝関連タンパク質に属する可能性が高いクローンはHEMBA1005732, MAMMA1000402の2クローンであった。DNA・RNA結合タンパク質に属する可能性が高いクローンはHEMBA1004596, OVARC1000148, PLACE1003334, THYR01001661の4クローンであった。タンパク質合成・輸送関連タンパク質に属する可能性が高いクローンはHEMBA1006284の1クローンであった。

【1096】残る2511クローンについては、現在のところ機能を推定できる情報の得られないクローンであった。これらクローンについては今後、機能が明らかに

2846

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2852

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2853

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2854

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2855

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2856

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2857

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2858

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【1120】THYRO1000270, THYRO1000279, THYRO10003 20, THYRO1000368, THYRO1000381, THYRO1000387, THYR
01000452, THYRO1000471, THYRO1000484, THYRO100050
2, THYRO1000505, THYRO1000558, THYRO1000596, THYRO

2859

1000625, THYR01000637, THYR01000676, THYR01000712, THYR01000715, THYR01000734, THYR01000777, THYR01000787, THYR01000793, THYR01000796, THYR01000805, THYR01000815, THYR01000843, THYR01000855, THYR01000865, THYR01000895, THYR01000916, THYR01000952, THYR01000988, THYR01001031, THYR01001062, THYR01001133, THYR01001142, THYR01001173, THYR01001213, THYR01001262, THYR01001321, THYR01001322, THYR01001363, THYR01001365, THYR01001403, THYR01001411, THYR01001426, THYR01001434, THYR01001480, THYR01001487, THYR01001559, THYR01001570, THYR01001584, THYR01001595, THYR01001602, THYR01001605, THYR01001637, THYR01001673, THYR01001706, THYR01001745, THYR01001746, THYR01001772, THYR01001793, THYR01001854, THYR01001895, THYR01001907, VESEN1000122, Y79AA1000065, Y79AA1000131, Y79AA1000202, Y79AA1000230, Y79AA1000355, Y79AA1000410, Y79AA1000480, Y79AA1000539, Y79AA1000574, Y79AA1000774, Y79AA1000802, Y79AA1000805, Y79AA1000824, Y79AA1000827, Y79AA1000850, Y79AA1000969, Y79AA1001041, Y79AA1001061, Y79AA1001068, Y79AA1001077, Y79AA1001078, Y79AA1001145, Y79AA1001167, Y79AA1001185, Y79AA1001216, Y79AA1001228, Y79AA1001281, Y79AA1001511, Y79AA1001541, Y79AA1001555, Y79AA1001585, Y79AA1001665, Y79AA1001696, Y79AA1001781, Y79AA1001805, Y79AA1002089, Y79AA1002115, Y79AA1002125, Y79AA1002220, Y79AA1002234, Y79AA1002298, Y79AA1002407, MAMMA1002215, MAMMA1002721, NT2RP2002070,

【1121】

【発明の効果】本発明により、5602にも及ぶポリヌクレオチドが提供された。全長cDNAの分離が進んでいないヒトにおいて、新規な全長cDNAを提供した意義は大きい。分泌蛋白質、膜蛋白質、シグナル伝達関連蛋白質、糖蛋白質関連蛋白質、転写関連蛋白質等は、多くの疾患に関連した蛋白質であることがわかっている。疾患に関連した遺伝子や蛋白質は、診断マーカー、発現や活性を制御する医薬品の開発、あるいは遺伝子治療のターゲットになるなど医薬品の開発等に有効である。中でも、分泌蛋白質をコードするcDNAは、蛋白質自身に医薬品としての有用性が期待できること、および多くの疾患に関連する遺伝子を含む可能性があることから、本発明によって提供されたこれらのcDNAは、産業上きわめて重要である。さらに、膜蛋白質、シグナル伝達関連蛋白質、転写関連蛋白質、あるいは疾患関連蛋白質といった蛋白質やそれをコードする遺伝子についても、疾患の指標となること等が期待できる。これらのcDNAも、産業上きわめて重要であり、コードする蛋白質の持つ活性の制御や、発現の制御を通じて疾患の治療効果をもたらすこと等が期待される。

【1122】 相同性検索結果データ1.

2860

5'末端クローン配列に対するSwissProt相同性検索結果データ

各データは、

クローン配列名、

トップヒットデータのDefinition、

P値: 比較配列の長さ (base): 相同性(%),

トップヒットデータのOrganism、

トップヒットデータのAccession No. の順に//で区切って記載した。

- 10 なお、相同性のスコアのP値が1より大であった場合はデータは示さない。ここでP値とは、配列間の類似性を統計的に起こりうる確率を考慮してスコアで示したもので、一般に値が小さいと類似性が高い(Altschul, S.F., Gish, W., Miller, W., Myers, E.W. & Lipman, D.J. (1990) "Basic local alignment search tool." J. Mol. Biol. 215:403-410; Gish, W. & States, D.J. (1993) "Identification of protein coding regions by database similarity search." Nature Genet. 3:266-272)。
- 20 F-HEMBA1000005//DNAJ PROTEIN HOMOLOG MTJ1.//1.8e-85:244:75//MUS MUSCULUS(MOUSE).//Q61712
F-HEMBA1000012//PROBABLE LEUCYL-TRNA SYNTHETASE (EC 6.1.1.4) (LEUCINE--TRNA LIGASE) (LEURS).//7.6e-57:231:53//CAENORHABDITIS ELEGANS.//Q09996
F-HEMBA1000020//TUBULIN BETA CHAIN.//1.0e-92:143:80//AJELLOWYCES CAPSULATA (HISTOPLASMA CAPSULATUM).//P41742
F-HEMBA1000030//CIRCUMSPOROZOITE PROTEIN PRECURSOR (CS).//0.021:136:33//PLASMODIUM KNOWLESII (STRAIN NURI).//P04922
F-HEMBA1000042//METALLOTHIONEIN 10-II (MT-10-II).//0.71:64:32//MYTILUS EDULIS (BLUE MUSSEL).//P80247
F-HEMBA1000046//PROTEIN Q300.//0.92:40:37//MUS MUSCULUS (MOUSE).//Q02722
F-HEMBA1000050//COMPETENCE PROTEIN S.//0.50:28:35//BACILLUS SUBTILIS.//P80355
F-HEMBA1000076//ATP SYNTHASE E CHAIN, MITOCHONDRIAL (EC 3.6.1.34).//0.86:41:41//HOMO SAPIENS (HUMAN).//P56385
- 40 F-HEMBA1000111
F-HEMBA1000129//UVSW PROTEIN (DAR PROTEIN).//0.023:68:33//BACTERIOPHAGE T4.//P20703
F-HEMBA1000141//YSY6 PROTEIN.//0.90:29:37//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38374
F-HEMBA1000150//!!!! ALU SUBFAMILY SP WARNING ENTRY!!!!//8.4e-16:47:70//HOMO SAPIENS (HUMAN).//P39193
F-HEMBA1000156//IMMEDIATE-EARLY PROTEIN.//8.1e-07:143:28//HERPESVIRUS SAIRI (STRAIN 11).//Q01042
- 50 F-HEMBA1000158//HYPOTHETICAL PROTEIN KIAA0192 (FRA

2861

CMENT). //7.9e-11:129:40//HOMO SAPIENS (HUMAN). //Q93074
 F-HEMBA1000168//INSULIN RECEPTOR SUBSTRATE-2 (IRS-2) (4PS). //0.00055:86:36//MUS MUSCULUS (MOUSE). //P81122
 F-HEMBA1000180//VP1 PROTEIN (U ORF PROTEIN). //0.22:73:28//CHIMPANZEE IMMUNODEFICIENCY VIRUS (SIV(CPZ)) (CIV). //P17286
 F-HEMBA1000185//RAS-1 PROTEIN. //5.1e-10:121:29//NEUROSPORA CRASSA. //P22126
 F-HEMBA1000193//PROLINE-RICH PEPTIDE P-B. //0.00078:56:41//HOMO SAPIENS (HUMAN). //P02814
 F-HEMBA1000201//PROLINE-RICH PROTEIN MP-2 PRECURSOR. //0.00061:49:42//MUS MUSCULUS (MOUSE). //P05142
 F-HEMBA1000213
 F-HEMBA1000216//HYPOXIA-INDUCIBLE FACTOR 1 ALPHA (HIF-1 ALPHA) (ARNT INTERACTING PROTEIN). //1.6e-59:115:53//MUS MUSCULUS (MOUSE). //Q61221
 F-HEMBA1000227//SUPPRESSOR PROTEIN SRP40. //0.00059:135:22//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32583
 F-HEMBA1000231//HYPOTHETICAL 60.7 KD PROTEIN C56F8.17C IN CHROMOSOME I. //0.024:60:38//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10264
 F-HEMBA1000243//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //0.0038:125:34//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1000244//HYPOTHETICAL 123.6 KD PROTEIN IN POR2-COX5B INTERGENIC REGION. //3.1e-17:149:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40480
 F-HEMBA1000251
 F-HEMBA1000264//PROBABLE E5 PROTEIN. //1.0:49:36//HUMAN PAPILLOMAVIRUS TYPE 58. //P26552
 F-HEMBA1000280//SHORT NEUROTOXIN 1 (TOXIN C-6). //0.98:58:31//NAJA NAJA KAOUTHIA (MONOCLED COBRA) (NAJA NAJA SIAMENSIS). //P14613
 F-HEMBA1000282//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!! //0.14:26:65//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1000288
 F-HEMBA1000290//HYPOTHETICAL 14 KD PROTEIN IN TVRI-6 REPETITIVE REGION. //3.8e-06:98:39//HOMO SAPIENS (HUMAN). //P10516
 F-HEMBA1000302
 F-HEMBA1000303//HYPOTHETICAL 104.4 KD PROTEIN F54G8.4 IN CHROMOSOME III. //1.3e-05:69:42//CAENORHABDITIS ELEGANS. //Q03601
 F-HEMBA1000304//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!! //0.021:18:83//HOMO SAPIENS (HUMAN). //P39194
 F-HEMBA1000307//MYOSIN HEAVY CHAIN, CLONE 203 (FRAGMENT). //7.1e-06:235:25//HYDRA ATTENUATA (HYDRA) (HYDRA VULGARIS). //P39922

2862

F-HEMBA1000327
 F-HEMBA1000333//SRP1 PROTEIN. //1.0:159:30//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10193
 F-HEMBA1000338//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!! //8.8e-26:36:83//HOMO SAPIENS (HUMAN). //P39193
 F-HEMBA1000351
 F-HEMBA1000355//BASIC PROLINE-RICH PEPTIDE P-E (IB-9). //0.99:22:50//HOMOSAPIENS (HUMAN). //P02811
 10 F-HEMBA1000356//IMMEDIATE-EARLY PROTEIN IE180. //0.11:82:36//PSEUDORABIESVIRUS (STRAIN INDIANA-FUNKHAUSER / BECKER) (PRV). //P11675
 F-HEMBA1000357//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!! //2.1e-35:105:74//HOMO SAPIENS (HUMAN). //P39192
 F-HEMBA1000366//HYPOTHETICAL TRANSCRIPTIONAL REGULATOR AF1627. //1.0:28:42//ARCHAEOGLOBUS FULGIDUS. //028646
 F-HEMBA1000369//PRESYNAPTIC DENSITY PROTEIN 95 (PSD-95). //0.013:140:26//HOMO SAPIENS (HUMAN). //P78352
 F-HEMBA1000376//RETROVIRUS-RELATED POLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); ENDONUCLEASE]. //6.8e-08:66:42//MUS MUSCULUS (MOUSE). //P11369
 F-HEMBA1000387//HYPOTHETICAL 63.2 KD PROTEIN C1F3.09 IN CHROMOSOME I. //1.5e-15:177:32//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10414
 F-HEMBA1000390//PARATHYMOSIN. //0.0071:61:29//HOMO SAPIENS (HUMAN). //P20962
 30 F-HEMBA1000392//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!! //2.1e-30:92:69//HOMO SAPIENS (HUMAN). //P39194
 F-HEMBA1000396//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //3.9e-23:64:57//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1000411
 F-HEMBA1000418
 F-HEMBA1000422//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!! //8.3e-10:90:53//HOMO SAPIENS (HUMAN). //P39188
 8 F-HEMBA1000428//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //3.1e-12:72:55//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1000434
 F-HEMBA1000442//GENE 11 PROTEIN. //1.0:28:46//SPIROPLASMA VIRUS SPV1-R8A2B. //P15902
 F-HEMBA1000456//26S PROTEASOME REGULATORY SUBUNIT MTS4 (19S REGULATORY CAP REGION OF 26S PROTEASE SUBUNIT 2). //0.077:118:28//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //P87048
 50 F-HEMBA1000459//HEME-REGULATED EUKARYOTIC INITIATION

2863

ON FACTOR EIF-2-ALPHA KINASE (EC 2.7.1.-) (HRI). // 4.8e-62:102:78//ORYZOLAGUS CUNICULUS (RABBIT). //P33279
 F-HEMBA1000460//LYSIS PROTEIN (E PROTEIN) (GPE). // 1.0:24:50//BACTERIOPHAGE ALPHA-3. //P31280
 F-HEMBA1000464
 F-HEMBA1000469//PILI PROTEIN. //1.0:27:44//PSEUDOMONAS AERUGINOSA. //P43502
 F-HEMBA1000488//ZINC FINGER PROTEIN 151 (MIZ-1 PROTEIN). //1.1e-07:90:38//HOMO SAPIENS (HUMAN). //Q13105
 F-HEMBA1000490//PLECTIN. //0.74:254:25//RATTUS NORVEGICUS (RAT). //P30427
 F-HEMBA1000491//RAS-RELATED PROTEIN M-RAS. //3.0e-14:100:36//RATTUS NORVEGICUS (RAT). //P97538
 F-HEMBA1000501//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//1.5e-20:81:54//HOMO SAPIENS (HUMAN). //P39194
 F-HEMBA1000504
 F-HEMBA1000505//NEURON-SPECIFIC X11 PROTEIN (FRAGMENT). //0.00028:128:32//HOMO SAPIENS (HUMAN). //Q02410
 F-HEMBA1000508//CHITIN SYNTHASE 3 (EC 2.4.1.16) (CHITIN-UDP ACETYL-GLUCOSAMINYL TRANSFERASE 3). //0.61:132:25//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P29465
 F-HEMBA1000518
 F-HEMBA1000519//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//1.8e-37:68:75//HOMO SAPIENS (HUMAN). //P39189
 F-HEMBA1000520//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!//5.2e-09:75:49//HOMO SAPIENS (HUMAN). //P39192
 F-HEMBA1000523//TESTIS-SPECIFIC PROTEIN PBS13. //1.5e-35:257:36//MUS MUSCULUS (MOUSE). //Q01755
 F-HEMBA1000531//HEAT SHOCK PROTEIN 70 B2. //1.6e-14:72:44//ANOPHELES ALBIMANUS (NEW WORLD MALARIA MOSQUITO). //P41827
 F-HEMBA1000534//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//9.7e-32:96:78//HOMO SAPIENS (HUMAN). //P39193
 F-HEMBA1000540//LANTIBIOTIC LACTICIN 481 PRECURSOR (LACTOCOCCIN DR). //1.0:12:75//LACTOCOCCUS LACTIS (SUBSP. LACTIS) (STREPTOCOCCUS LACTIS). //P36499
 F-HEMBA1000542//SPERM MITOCHONDRIAL CAPSULE SELENO PROTEIN (MCS). //0.0089:79:31//MUS MUSCULUS (MOUSE). //P15265
 F-HEMBA1000545//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //9.0e-83:256:66//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1000555//TRANSLATION INITIATION FACTOR IF-

2864

2. //3.6e-06:252:22//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P39730
 F-HEMBA1000557
 F-HEMBA1000561//ZINC FINGER PROTEIN 81 (FRAGMENT). //9.1e-18:200:28//HOMO SAPIENS (HUMAN). //P51508
 F-HEMBA1000563
 F-HEMBA1000568
 F-HEMBA1000569//GPI-ANCHORED PROTEIN P137. //1.0e-40:137:54//HOMO SAPIENS (HUMAN). //Q14444
 F-HEMBA1000575
 F-HEMBA1000588
 F-HEMBA1000591//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//3.1e-17:41:92//HOMO SAPIENS (HUMAN). //P39194
 F-HEMBA1000592//CENTROMERIC PROTEIN E (CENP-E PROTEIN). //0.18:128:23//HOMO SAPIENS (HUMAN). //Q02224
 F-HEMBA1000594//HYPOTHETICAL 29.3 KD PROTEIN B028.0.6 IN CHROMOSOME III. //0.93:24:54//CAENORHABDITIS ELEGANS. //P41997
 F-HEMBA1000604//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//0.00010:49:55//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1000608//HYPOTHETICAL PROTEIN KIAA0411 (FRAGMENT). //1.8e-55:179:61//HOMO SAPIENS (HUMAN). //Q43295
 F-HEMBA1000622//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.1e-21:94:62//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1000636//SULFATED SURFACE GLYCOPROTEIN 185 (SSG 185). //0.34:73:36//VOLVOX CARTERI. //P21997
 F-HEMBA1000637//BASIC PROLINE-RICH PEPTIDE IB-1. //0.0057:76:38//HOMO SAPIENS (HUMAN). //P04281
 F-HEMBA1000655
 F-HEMBA1000657//ZINC FINGER PROTEIN GCS1. //1.5e-07:66:37//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P35197
 F-HEMBA1000662//METALLOTHIONEIN-II (MT-II). //0.79:33:39//CRICETULUS GRISEUS (CHINESE HAMSTER). //P02799
 F-HEMBA1000673//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//3.1e-17:86:59//HOMO SAPIENS (HUMAN). //P39193
 F-HEMBA1000682//RETROVIRUS-RELATED POLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); ENDONUCLEASE]. //3.0e-13:45:44//MUS MUSCULUS (MOUSE). //P11369
 F-HEMBA1000686//HYPOTHETICAL 48.0 KD PROTEIN C1B3.08 IN CHROMOSOME I. //4.5e-07:79:34//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q13873
 F-HEMBA1000702

2865

F-HEMBA1000705//PROTEIN Q300.//0.80:25:44//MUS MUS
CULUS (MOUSE).//Q02722
F-HEMBA1000719//MYOSIN IC HEAVY CHAIN.//0.0026:11
5:44//ACANTHAMOEBA CASTELLANII (AMOEBA).//P10569
F-HEMBA1000722
F-HEMBA1000726//!!!! ALU SUBFAMILY SB2 WARNING ENT
RY !!!!!/7.4e-32:83:77//HOMO SAPIENS (HUMAN).//P39
191
F-HEMBA1000727//ZINC FINGER PROTEIN CTH2 (YTIS11 P
ROTEIN).//0.73:26:46//SACCHAROMYCES CEREVISIAE (BA 10
KER'S YEAST).//P47977
F-HEMBA1000747
F-HEMBA1000749//HYPOTHETICAL PROTEIN HI1484.//1.0:
42:35//HAEMOPHILUS INFLUENZAE.//P44211
F-HEMBA1000752//RETROVIRUS-RELATED ENV POLYPROTEI
N.//1.0e-08:84:39//HOMOSAPIENS (HUMAN).//P10267
F-HEMBA1000769
F-HEMBA1000773//PAIRED BOX PROTEIN PAX-4.//1.0:10
7:33//HOMO SAPIENS (HUMAN).//043316
F-HEMBA1000774//!!!! ALU SUBFAMILY J WARNING ENTRY 20
!!!!/1.3e-23:92:63//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1000791
F-HEMBA1000817//PROLACTIN RECEPTOR PRECURSOR (PRL-
R).//0.079:87:29//CERVUS ELAPHUS (RED DEER).//Q282
35
F-HEMBA1000822
F-HEMBA1000827//HYPOTHETICAL 8.4 KD PROTEIN.//0.9
8:48:39//VACCINIA VIRUS(STRAIN COPENHAGEN).//P2054
6
F-HEMBA1000843//HYPOTHETICAL 7.3 KD PROTEIN D1044.
5 IN CHROMOSOME III.//0.92:46:34//CAENORHABDITIS E
LEGANS.//P41953
F-HEMBA1000851//HOMEBOX PROTEIN GBX-2 (GASTRULATI
ON AND BRAIN-SPECIFICHOMEBOX PROTEIN 2).//0.048:3
9:51//HOMO SAPIENS (HUMAN).//P52951
F-HEMBA1000852//ARYLSULFATASE D PRECURSOR (EC 3.1.
6.-) (ASD).//4.0e-24:29:100//HOMO SAPIENS (HUMAN).
//P51689
F-HEMBA1000867
F-HEMBA1000869//PROBABLE E5 PROTEIN.//0.99:70:27//
HUMAN PAPILLOMAVIRUS TYPE 18.//P06792
F-HEMBA1000870//MYOTOXIN 3 PRECURSOR (CROTAMINE
3).//0.79:43:32//CROTALUS DURISSUS TERRIFICUS (SOU
TH AMERICAN RATTLESNAKE).//P24333
F-HEMBA1000872//GAR2 PROTEIN.//0.89:70:31//SCHIZO
SACCHAROMYCES POMBE (FISSION YEAST).//P41891
F-HEMBA1000876//DEFENSIN.//0.89:34:38//ALLOWYRINA
DICHOTOMA.//Q10745
F-HEMBA1000908//DNA-BINDING P52/P100 COMPLEX, 100 50

2866

KD SUBUNIT (FRAGMENTS).//0.69:43:37//HOMO SAPIENS
(HUMAN).//P30808
F-HEMBA1000910//MELANOMA-ASSOCIATED ANTIGEN B3 (MA
GE-B3 ANTIGEN).//5.1e-08:44:38//HOMO SAPIENS (HUMA
N).//015480
F-HEMBA1000918//60S RIBOSOMAL PROTEIN L37-A (YL35)
(FRAGMENT).//1.0:19:52//SCHIZOSACCHAROMYCES POMBE
(FISSION YEAST).//P22667
F-HEMBA1000919//69 KD PARAFLAGELLAR ROD PROTEIN (6
9 KD PFR PROTEIN) (PFR-A/PFR-B).//0.29:116:30//TRY
PANOSOMA BRUCEI BRUCEI.//P22225
F-HEMBA1000934
F-HEMBA1000942//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/0.85:27:59//HOMO SAPIENS (HUMAN).//P39188
F-HEMBA1000943
F-HEMBA1000946//STO-2 PROTEIN.//0.82:82:30//CAENOR
HABDITIS ELEGANS.//Q19958
F-HEMBA1000960//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!/0.0097:29:72//HOMO SAPIENS (HUMAN).//P3919
2
F-HEMBA1000968//METALLOTHIONEIN 20-III ISOFORMS A
AND B (WT-20-IIIA AND WT-20-IIIB).//0.047:45:37//MY
TILUS EDULIS (BLUE MUSSEL).//P80253
F-HEMBA1000971//HYPOTHETICAL BHLF1 PROTEIN.//0.03
8:172:31//EPSTEIN-BARRVIRUS (STRAIN B95-8) (HUMAN
HERPESVIRUS 4).//P03181
F-HEMBA1000972
F-HEMBA1000974//HYPOTHETICAL PROTEIN MC441.//0.98:
66:28//MYCOPLASMA GENITALIUM.//P47679
30 F-HEMBA1000975//COLLAGEN ALPHA 2(VIII) CHAIN (ENDO
THELIAL COLLAGEN) (FRAGMENT).//0.028:57:36//HOMO S
APIENS (HUMAN).//P25067
F-HEMBA1000985
F-HEMBA1000986//SUBMANDIBULAR GLAND SECRETORY GLX-
RICH PROTEIN CB PRECURSOR (GRP-CB) (CONTIGUOUS REP
EAT POLYPEPTIDE) (CRP).//0.13:91:34//RATTUSNORVEGI
CUS (RAT).//P08462
F-HEMBA1000991//HYPOTHETICAL 46.2 KD TRP-ASP REPEA
TS CONTAINING PROTEIN D2013.2 IN CHROMOSOME II.//5.
40 6e-05:37:45//CAENORHABDITIS ELEGANS.//Q18964
F-HEMBA1001007//HYPOTHETICAL PROTEIN KIAA0179.//0.
27:72:41//HOMO SAPIENS (HUMAN).//Q14684
F-HEMBA1001008//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/2.4e-25:61:70//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBA1001009//CUTICLE COLLAGEN 34.//0.044:214:29
//CAENORHABDITIS ELEGANS.//P34687
F-HEMBA1001017//SYNDECAN-3 PRECURSOR (N-SYNDECAN)
(NEUROGLYCAN).//5.0e-85:191:84//RATTUS NORVEGICUS
(RAT).//P33671

F-HEMBA1001019
 F-HEMBA1001020//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/6.7e-24:49:73//HOMO SAPIENS (HUMAN).//P3918
 8
 F-HEMBA1001022
 F-HEMBA1001024//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG.//3.0e-11:61:59//HOMO SAPIENS (HUMAN).//P08547
 F-HEMBA1001026//HYPOTHETICAL PROTEIN BB0073.//0.9
 4:63:34//BORRELIA BURGDORFERI (LYME DISEASE SPIROCH
 ETE).//051100
 F-HEMBA1001043//INVOLUCRIN.//0.0036:238:25//SAGUIN
 US OEDIPUS (COTTON-TOPTAMARIN).//P24712
 F-HEMBA1001051//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/3.3e-32:95:75//HOMO SAPIENS (HUMAN).//P391
 89
 F-HEMBA1001052//CURROMYCIN RESISTANCE PROTEIN.//1.
 0:31:38//STREPTOMYCESHYGROSCOPICUS.//P16961
 F-HEMBA1001059//N-ACETYLGALACTOSAMINE-6-SULFATASE
 PRECURSOR (EC 3.1.6.4) (N- ACETYLGALACTOSAMINE-6-SU
 LFATE SULFATASE) (GALACTOSE-6-SULFATE SULFATASE)
 (GALNAC6S SULFATASE) (CHONDROITINSULFATASE) (CHOND
 ROITINASE).//3.2e-132:249:94//HOMO SAPIENS (HUMA
 N).//P34059
 F-HEMBA1001060
 F-HEMBA1001071//PROCOLLAGEN ALPHA 1(III) CHAIN PRE
 CURSOR.//8.3e-23:51:96//HOMO SAPIENS (HUMAN).//P02
 461
 F-HEMBA1001077//AUTOIMMUNE REGULATOR (APECED PROTE
 IN).//3.4e-06:37:56//HOMO SAPIENS (HUMAN).//043918
 F-HEMBA1001080//INFECTED CELL PROTEIN ICP34.5 (NEU
 30 ROVIRULENCE FACTOR ICP34.5).//0.0012:70:38//HERPES
 SIMPLEX VIRUS (TYPE 1 / STRAIN MGH-10).//P37319
 F-HEMBA1001085//SERINE/THREONINE PROTEIN PHOSPHATA
 SE 5 (EC 3.1.3.16) (PP5) (PROTEIN PHOSPHATASE T)
 (PPT) (FRAGMENT).//0.00018:76:32//MUS MUSCULUS (MO
 USE).//Q60676
 F-HEMBA1001088//PINCH PROTEIN (PARTICULARY INTERES
 TING NEW CYS-HIS PROTEIN).//3.5e-50:176:57//HOMO S
 APIENS (HUMAN).//P48059
 F-HEMBA1001094
 F-HEMBA1001099//LIGHT-HARVESTING PROTEIN B800/850/
 890, ALPHA-2 CHAIN (EHA-ALPHA-2) (ANTENNA PIGMENT
 PROTEIN, ALPHA-2 CHAIN) (FRAGMENT).//1.0:15:60//EC
 TOTHIORHODOSPIRA HALOPHILA.//P80101
 F-HEMBA1001109//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/6.7e-37:102:82//HOMO SAPIENS (HUMAN).//P39
 189
 F-HEMBA1001121//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG.//0.036:49:46//HOMOSAPIENS (HUMAN).//P08547
 F-HEMBA1001122

F-HEMBA1001123
 F-HEMBA1001133//HYPOTHETICAL 9.4 KD PROTEIN (ORF
 2).//0.86:29:41//FELINEIMMUNODEFICIENCY VIRUS (ISO
 LATE SAN DIEGO) (FIV), AND FELINE IMMUNODEFICIENCY
 VIRUS (ISOLATE PETALUMA) (FIV).//P19033
 F-HEMBA1001137//ZINC FINGER PROTEIN 84 (ZINC FINGE
 R PROTEIN HPF2).//2.0e-22:103:52//HOMO SAPIENS (HU
 MAN).//P51523
 F-HEMBA1001140//COLLAGEN ALPHA 4(IV) CHAIN PRECURS
 10 OR.//0.032:94:36//HOMOSAPIENS (HUMAN).//P53420
 F-HEMBA1001172
 F-HEMBA1001174//ADP-RIBOSYLATION FACTOR-LIKE PROTE
 IN 5.//2.9e-78:179:79//RATTUS NORVEGICUS (RAT).//P
 51646
 F-HEMBA1001197//MAJOR PRION PROTEIN PRECURSOR (PR
 P) (PRP27-30) (PRP33-35C) (FRAGMENT).//0.051:96:32
 //CERCOCEBUS ATERRIMUS, AND MACACA SYLVANUS (BARBA
 RY APE).//Q95145
 F-HEMBA1001208
 20 F-HEMBA1001213
 F-HEMBA1001226//PROTEASOME COMPONENT C8 (EC 3.4.9
 9.46) (MACROPAIN SUBUNIT C8) (MULTICATALYTIC ENDOP
 EPTIDASE COMPLEX SUBUNIT C8).//1.5e-08:24:91//HOMO
 SAPIENS (HUMAN).//P25788
 F-HEMBA1001235//FIBRONECTIN (FN) (FRAGMENT).//0.7
 6:50:38//ORYCTOLAGUS CUNICULUS (RABBIT).//Q28749
 F-HEMBA1001247//SULFATED SURFACE GLYCOPROTEIN 185
 (SSG 185).//0.00052:16:81//VOLVOX CARTERI.//P21997
 F-HEMBA1001257//2-ARYLPROPIONYL-COA EPIMERASE (EC
 5.-.-.-).//1.6e-68:178:77//RATTUS NORVEGICUS (RA
 30 T).//P70473
 F-HEMBA1001265//MANNAN ENDO-1,4-BETA-MANNOSIDASE A
 PRECURSOR (EC 3.2.1.78) (BETA- MANNANASE A) (1,4-
 BETA-D-MANNAN MANNANOHYDROLASE A).//0.67:23:60//PI
 ROMYCES SP.//P55296
 F-HEMBA1001281//HYPOTHETICAL 8.9 KD PROTEIN YCF34
 (ORF76).//0.83:48:35//PORPHYRA PURPUREA.//P51229
 F-HEMBA1001286//COMPLEMENT DECAY-ACCELERATING FACT
 OR PRECURSOR.//1.3e-07:185:29//CAVIA PORCELLUS (GU
 40 INEA PIG).//Q60401
 F-HEMBA1001289//METABOTROPIC GLUTAMATE RECEPTOR 3
 PRECURSOR.//0.00018:159:30//RATTUS NORVEGICUS (RA
 T).//P31422
 F-HEMBA1001294
 F-HEMBA1001299//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!/5.3e-07:27:77//HOMO SAPIENS (HUMAN).//P391
 95
 F-HEMBA1001302//45 KD CALCIUM-BINDING PROTEIN PREC
 ursor (STROMAL CELL-DERIVED FACTOR 4) (SDF-4).//3.
 50 3e-61:150:76//MUS MUSCULUS (MOUSE).//Q61112

- F-HEMBA1001303
 F-HEMBA1001310//HYPOTHETICAL PROTEIN KIAA0161.//2.
 7e-10:170:27//HOMO SAPIENS (HUMAN).//P50876
 F-HEMBA1001319
 F-HEMBA1001323
 F-HEMBA1001326//HYPOTHETICAL 55.1 KD PROTEIN IN FA
 B1-PES4 INTERGENIC REGION.//1.1e-39:144:38//SACCHA
 ROMYCES CEREVISIAE (BAKER'S YEAST).//P43601
 F-HEMBA1001327
 F-HEMBA1001330
 F-HEMBA1001351//VESICLE-ASSOCIATED MEMBRANE PROTEI
 N/SYNAPOBREVIN BINDING PROTEIN (VAP-33).//1.9e-3
 7:155:46//APLYSIA CALIFORNICA (CALIFORNIA SEAHAR
 E).//Q16943
 F-HEMBA1001361//RUBREDOXIN (RD).//0.95:44:29//ALCA
 LIGENES EUTROPHUS.//P31912
 F-HEMBA1001375//AEROLYSIN REGULATORY PROTEIN.//0.0
 13:45:33//AEROMONAS SOBRIA.//P09165
 F-HEMBA1001377//SPERM PROTAMINE P1.//1.0:22:40//PL
 ANIGALE MACULATA SINUALIS (COMMON PLANIGALE).//018
 746
 F-HEMBA1001383//TAT PROTEIN (TRANSACTIVATING REGUL
 ATORY PROTEIN).//0.60:37:29//HUMAN IMMUNODEFICIENC
 Y VIRUS TYPE 1 (Z2/CDC-Z34 ISOLATE) (HIV-1).//P125
 06
 F-HEMBA1001387//GTP-BINDING PROTEIN TC10.//6.6e-4
 3:83:92//HOMO SAPIENS (HUMAN).//P17081
 F-HEMBA1001388//HYPOTHETICAL PROTEIN KIAA0136 (FRA
 GMENT).//0.00088:46:45//HOMO SAPIENS (HUMAN).//Q14
 149
 F-HEMBA1001391
 F-HEMBA1001398//CLOACIN (EC 3.1.-.-) (RIBONUCLEAS
 E).//1.0:59:37//ESCHERICHIA COLI.//P00645
 F-HEMBA1001405//SMALL PROLINE RICH PROTEIN II (SPR
 -II) (CLONE 930).//0.25:41:34//HOMO SAPIENS (HUMA
 N).//P22531
 F-HEMBA1001407//SALIVARY PROLINE-RICH PROTEIN PREC
 URSOR (CLONES CP3, CP4AND CP5) [CONTAINS: BASIC PE
 PTIDE IB-6; PEPTIDE P-H].//4.0e-09:129:40//HOMO SA
 PIENS (HUMAN).//P04280
 F-HEMBA1001411//HYPOTHETICAL 34.9 KD PROTEIN IN CY
 SJ-ENO INTERGENIC REGION (0313).//0.95:88:31//ESCH
 ERICHIA COLI.//P55140
 F-HEMBA1001413//SOX-12 PROTEIN (FRAGMENT).//0.95:4
 6:32//MUS MUSCULUS (MOUSE).//Q04890
 F-HEMBA1001415//HISTONE H5.//0.43:95:29//GALLUS GA
 LLUS (CHICKEN).//P02259
 F-HEMBA1001432//LANTIBIOTIC NISIN A PRECURSOR.//0.
 77:46:32//LACTOCOCCUS LACTIS (SUBSP. LACTIS) (STREP
 TOCOCUS LACTIS).//P13068
- F-HEMBA1001433//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG.//4.8e-09:132:31//NYCTICEBUS COUCANG (SLOW LORI
 S).//P08548
 F-HEMBA1001435//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!//1.2e-31:84:77//HOMO SAPIENS (HUMAN).//P391
 89
 F-HEMBA1001442
 F-HEMBA1001446//ANTIFREEZE PEPTIDE 4 PRECURSOR.//
 0.71:41:39//PSEUDOPLEURONECTA AMERICANUS (WINTER F
 LOUNDER).//P02734
 F-HEMBA1001450//PROLINE-RICH PROTEIN LAS17.//0.13:
 127:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//
 Q12446
 F-HEMBA1001454//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
 4) (AGL).//0.57:38:47//HANSENULA WINGEI (YEAST).//
 P48882
 F-HEMBA1001455//CHEMOTAXIS PROTEIN CHEA (EC 2.7.3.
 -).//0.98:124:25//BORRELIA BURGDORFERI (LYME DISEA
 SE SPIROCHETE).//Q44737
 F-HEMBA1001463//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//2.8e-32:62:67//HOMO SAPIENS (HUMAN).//P391
 94
 F-HEMBA1001476//NUCLEOPORIN NUP159 (NUCLEAR PORE P
 ROTEIN NUP159).//6.8e-09:252:36//SACCHAROMYCES CER
 EVISIAE (BAKER'S YEAST).//P40477
 F-HEMBA1001478
 F-HEMBA1001497//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//1.2e-33:105:72//HOMO SAPIENS (HUMAN).//P39
 194
 F-HEMBA1001510//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!//8.3e-37:54:81//HOMO SAPIENS (HUMAN).//P391
 89
 F-HEMBA1001515//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG.//2.0e-63:223:57//HOMO SAPIENS (HUMAN).//P08547
 F-HEMBA1001517
 F-HEMBA1001522//TROPOMYOSIN ALPHA CHAIN, SMOOTH MU
 SCLE.//0.78:150:22//COTURNIX COTURNIX JAPONICA (JA
 PANESE QUAIL).//P49437
 F-HEMBA1001526//PERIPLASMIC [FE] HYDROGENASE 1 (EC
 1.18.99.1).//1.6e-06:130:29//CLOSTRIDIUM PASTEUR
 ANUM.//P29166
 F-HEMBA1001533//PROBABLE E5A PROTEIN.//0.73:35:37/
 /HUMAN PAPILOMAVIRUS TYPE 6A.//Q84296
 F-HEMBA1001557//HYPOTHETICAL 17.1 KD PROTEIN IN PU
 R5 3' REGION.//1.5e-07:99:36//SACCHAROMYCES CEREVIS
 IAE (BAKER'S YEAST).//P38898
 F-HEMBA1001566//HYPOTHETICAL PROTEIN BB0692.//0.9
 1:27:44//BORRELIA BURGDORFERI (LYME DISEASE SPIROC
 HETE).//051635
 F-HEMBA1001569//SYNAPOBREVIN 2 (VESICLE ASSOCIATE

2871

D MEMBRANE PROTEIN 2) (VAMP-2). //2.2e-50:110:95//HOMO SAPIENS (HUMAN), AND BOS TAURUS (BOVINE). //P19065
 F-HEMBA1001570//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//3.3e-33:107:72//HOMO SAPIENS (HUMAN). //P39195
 F-HEMBA1001579//RING CANAL PROTEIN (KELCH PROTEIN). //1.2e-14:111:39//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q04652
 F-HEMBA1001581
 F-HEMBA1001585
 F-HEMBA1001589//PROBABLE DNA-BINDING PROTEIN (AGNO PROTEIN). //0.98:51:33//HUMAN ADENOVIRUS TYPE 2. //P03263
 F-HEMBA1001595//SEPTIN 2 HOMOLOG (FRAGMENT). //3.0e-124:274:85//HOMO SAPIENS (HUMAN). //Q14141
 F-HEMBA1001608//RENAL SODIUM/DICARBOXYLATE COTRANSPORTER (NA(+)/DICARBOXYLATE COTRANSPORTER). //0.99:28:39//ORYZOLAGUS CUNICULUS (RABBIT). //Q28615
 F-HEMBA1001620//MYO-INOSITOL-1-PHOSPHATE SYNTHASE (EC 5.5.1.4) (IPS). //4.3e-45:222:46//SPIRODELA POLYRRHIZA. //P42803
 F-HEMBA1001635//FIBRILLARIN. //0.10:72:38//CAENORHABDITIS ELEGANS. //Q22053
 F-HEMBA1001636//PAIRED BOX PROTEIN PAX-8, ISOFORMS 8C/8D. //0.75:38:47//HOMO SAPIENS (HUMAN). //Q09155
 F-HEMBA1001640//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//4.7e-06:80:41//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1001647//TRANSCRIPTION INITIATION FACTOR TFIID 135 KD SUBUNIT (TAFII-135) (TAFII135) (TAFII-130) (TAFII130). //0.075:165:32//HOMO SAPIENS (HUMAN). //Q00268
 F-HEMBA1001651//GOLGIN-95. //6.8e-05:141:24//HOMO SAPIENS (HUMAN). //Q08379
 F-HEMBA1001655//PROLINE-RICH PROTEIN LAS17. //0.19:97:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q12446
 F-HEMBA1001658//TETRAHYDROMETHANOPTERIN S-METHYLTRANSFERASE 12 KD SUBUNIT (EC 2.1.1.86) (N5-METHYLTRANSFERASE 12 KD SUBUNIT). //1.0:29:44//METHANOBACTERIUM THERMAUTOTROPHICUM (STRAIN MARBURG / DSM 2133). //Q50773
 F-HEMBA1001661//CELLULOSE COMPLEMENTING PROTEIN. //0.35:87:33//ACETOBACTER XYLINUM (ACETOBACTER PASTERURIANUS). //P37697
 F-HEMBA1001672//CIRCUMSPOROZOITE PROTEIN PRECURSOR (CS). //2.7e-10:216:35//PLASMODIUM CYNOMOLGI (STRAIN BEROK). //P08672

2872

F-HEMBA1001675//NODULIN 20 PRECURSOR (N-20). //0.98:36:44//GLYCINE MAX (SOYBEAN). //P08960
 F-HEMBA1001678//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//8.2e-13:62:64//HOMO SAPIENS (HUMAN). //P39195
 F-HEMBA1001681//HYPOTHETICAL 41.5 KD PROTEIN IN P6.5-VP48 INTERGENIC REGION (P40) (ORF3) (ORF102). //1.0:51:39//ORGANIA PSEUDOTSUGATA MULTICAPSID POLYHEROSIS VIRUS (OPMNPV). //P24653
 10 F-HEMBA1001702//ATP SYNTHASE A CHAIN (EC 3.6.1.34) (PROTEIN 6). //0.017:54:37//TRYPANOSOMA BRUCEI BRUCEI. //P24499
 F-HEMBA1001709//HYPOTHETICAL 21.2 KD PROTEIN IN TO R2-MNN4 INTERGENIC REGION. //0.59:109:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P36042
 F-HEMBA1001711
 F-HEMBA1001712//HYPOTHETICAL 6.9 KD PROTEIN IN 100 KD PROTEIN REGION. //0.54:44:34//HUMAN ADENOVIRUS TYPE 41. //P23690
 20 F-HEMBA1001714//ATPASE INHIBITOR, MITOCHONDRIAL PRECURSOR. //1.2e-19:60:75//RATTUS NORVEGICUS (RAT). //Q03344
 F-HEMBA1001718//HYPOTHETICAL PROTEIN UL63. //1.0:54:37//HUMAN CYTOMEGALOVIRUS (STRAIN AD169). //P16820
 F-HEMBA1001723//HYPOTHETICAL 34.0 KD TRP-ASP REPEATS CONTAINING PROTEININ SIS1-MRPL2 INTERGENIC REGION. //5.1e-26:90:53//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P41318
 30 F-HEMBA1001731//HYPOTHETICAL 16.6 KD PROTEIN. //0.71:49:32//AVIAN INFECTIOUS BURSAL DISEASE VIRUS (STRAIN 52/70) (IBDV). //P25221
 F-HEMBA1001734
 F-HEMBA1001744//SCY1 PROTEIN. //2.1e-11:182:29//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53009
 F-HEMBA1001745//HYPOTHETICAL 11.6 KD PROTEIN IN NUT1-AR02 INTERGENIC REGION PRECURSOR. //1.0:36:38//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53116
 F-HEMBA1001746//PROTEIN-EXPORT MEMBRANE PROTEIN SEC HOMOLOG. //0.94:48:35//MYCOBACTERIUM LEPRAE. //P38388
 F-HEMBA1001761
 F-HEMBA1001781//ZINC FINGER PROTEIN 19 (ZINC FINGER PROTEIN KOX12) (FRAGMENT). //0.028:47:40//HOMO SAPIENS (HUMAN). //P17023
 F-HEMBA1001784//HYPOTHETICAL 6.1 KD PROTEIN CO3B1.10 IN CHROMOSOME X. //0.00068:32:46//CAENORHABDITIS ELEGANS. //Q11116
 F-HEMBA1001791//METALLOTHIONEIN (MT). //1.0:34:35//PLEURONECTES PLATESSA (PLAICE). //P07216
 50

- F-HEMBA1001800//ZINC FINGER PROTEIN MFG-1 (ZINC FINGER PROTEIN 58) (FRAGMENT).//1.5e-14:60:48//MUS MUSCULUS (MOUSE).//P16372
- F-HEMBA1001803
- F-HEMBA1001804//GLYCINE-RICH CELL WALL STRUCTURAL PROTEIN 1 PRECURSOR.//9.3e-17:56:57//ORYZA SATIVA (RICE).//P25074
- F-HEMBA1001808//PARANEOPLASTIC ENCEPHALOMYELITIS ANTIGEN HUD HOMOLOG (HU-ANTIGEN D).//0.75:97:31//RATTUS NORVEGICUS (RAT).//009032
- F-HEMBA1001809//IMMEDIATE-EARLY PROTEIN IE180.//4.5e-11:206:36//PSEUDORABIES VIRUS (STRAIN INDIANA-FUNKHAUSER / BECKER) (PRV).//P11675
- F-HEMBA1001815//60S RIBOSOMAL PROTEIN L37-B (YL27) (FRAGMENT).//0.34:30:30//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//P05733
- F-HEMBA1001819//ZINC FINGER PROTEIN 135.//2.6e-10:2:262:66//HOMO SAPIENS (HUMAN).//P52742
- F-HEMBA1001820
- F-HEMBA1001822//EPIDERMAL GROWTH FACTOR RECEPTOR SUBSTRATE SUBSTRATE 15 (PROTEIN EPS15).//1.2e-18:25:1:33//MUS MUSCULUS (MOUSE).//P42567
- F-HEMBA1001824//KERATIN, ULTRA HIGH-SULFUR MATRIX PROTEIN (UHS KERATIN).//4.7e-11:124:37//OVIS ARIES (SHEEP).//P26372
- F-HEMBA1001835
- F-HEMBA1001844//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!//4.3e-14:36:63//HOMO SAPIENS (HUMAN).//P39195
- F-HEMBA1001847//ZINC FINGER PROTEIN 29 (ZFP-29).//2.7e-36:135:51//MUS MUSCULUS (MOUSE).//Q07230
- F-HEMBA1001861
- F-HEMBA1001864//HEAT-STABLE ENTEROTOXIN A3/A4 PRECURSOR (STA3/STA4) (ST-IB) (ST-H).//1.0:31:38//ESCHERICHIA COLI.//P07965
- F-HEMBA1001866//UDP-GLUCOSE: GLYCOPROTEIN GLUCOSYLTRANSFERASE PRECURSOR (EC 2.4.1.-) (DUGT).//9.7e-4:2:234:41//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q09332
- F-HEMBA1001869//HYPOTHETICAL 94.9 KD PROTEIN C22E1 2.11C IN CHROMOSOME I.//5.3e-13:65:47//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q10362
- F-HEMBA1001888//HYPOTHETICAL 11.4 KD PROTEIN (ORF 1).//0.85:62:37//STREPTOMYCES FRADIAE.//P26800
- F-HEMBA1001896//DIMETHYLGLYCINE DEHYDROGENASE PRECURSOR (EC 1.5.99.2) (ME2GLYDH).//9.8e-20:250:29//RATTUS NORVEGICUS (RAT).//Q63342
- F-HEMBA1001910//EUKARYOTIC TRANSLATION INITIATION FACTOR 4E (EIF-4E) (EIF4E) (MRNA CAP-BINDING PROTEIN) (EIF-4F 25 KD SUBUNIT).//0.94:44:38//CAENORHABDITIS ELEGANS.//061955
- F-HEMBA1001912//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!//8.7e-07:53:62//HOMO SAPIENS (HUMAN).//P39188
- F-HEMBA1001913//GCN20 PROTEIN.//1.8e-21:68:60//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P43535
- F-HEMBA1001915//KLEE PROTEIN (KCRB3 PROTEIN).//0.94:64:21//ESCHERICHIA COLI.//Q52280
- F-HEMBA1001918
- F-HEMBA1001921
- F-HEMBA1001939//CHLOROPLAST 50S RIBOSOMAL PROTEIN L24.//1.0:47:31//ODONTELLA SINENSIS.//P49560
- F-HEMBA1001940//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!//0.0017:31:77//HOMO SAPIENS (HUMAN).//P39188
- F-HEMBA1001942//HIBERNATION-ASSOCIATED PLASMA PROTEIN HP-27 PRECURSOR (HIBERNATOR-SPECIFIC BLOOD COMPLEX, 27 KD SUBUNIT).//1.0:77:28//TAMIAS ASIATICUS (CHIPMUNK).//Q06577
- F-HEMBA1001945//HYPOTHETICAL 4.6 KD PROTEIN IN GP4 7-ACT INTERGENIC REGION (ORF E).//1.0:35:37//BACTERIOPHAGE T4.//P32269
- F-HEMBA1001950//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3 4) (A6L).//0.43:18:44//DROSOPHILA YAKUBA (FRUIT FLY).//P03933
- F-HEMBA1001960//HOMEODOMAIN PROTEIN HOX-C5 (HOX-3D) (CP11).//0.17:12:66//HOMO SAPIENS (HUMAN).//Q00444
- F-HEMBA1001962//HYPOTHETICAL 9.0 KD PROTEIN IN ADH 4 5' REGION.//1.0:30:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53056
- F-HEMBA1001964
- F-HEMBA1001967//HYPOTHETICAL PROTEIN UL61.//0.027:111:36//HUMAN CYTOMEGALOVIRUS (STRAIN AD169).//P16818
- F-HEMBA1001979
- F-HEMBA1001987//HYPOTHETICAL 11.2 KD PROTEIN (ORF 17).//1.0:83:32//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPMNPV).//O10356
- F-HEMBA1001991//NEUROTOXIN 1 (TOXIN ATX-I).//0.99:31:45//ANEMONIA SULCATA (SNAKE-LOCKS SEA ANEMONE).//P01533
- F-HEMBA1002003//GLYCERALDEHYDE 3-PHOSPHATE DEHYDROGENASE, TESTIS-SPECIFIC (EC 1.2.1.12) (GAPDH).//5.5e-07:109:32//MUS MUSCULUS (MOUSE).//Q64467
- F-HEMBA1002008
- F-HEMBA1002018//EC PROTEIN HOMOLOG 2 (FRAGMENT).//0.83:66:33//ARABIDOPSIS THALIANA (MOUSE-EAR CRUCIFER).//Q42377
- F-HEMBA1002022//INSULIN.//1.0:59:32//SQUALUS ACANTHIAS (SPINY DOGFISH).//P12704
- F-HEMBA1002035//MONOCYTIC LEUKEMIA ZINC FINGER PRO

2875

2876

TEIN. //8.3e-15:64:40//HOMO SAPIENS (HUMAN). //Q92794

F-HEMBA1002039//HYPOTHETICAL PROLINE-RICH PROTEIN KIAA0269. //0.0070:70:40//HOMO SAPIENS (HUMAN). //Q92558

F-HEMBA1002049//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.5e-07:37:75//HOMO SAPIENS (HUMAN). //P39188

F-HEMBA1002084

F-HEMBA1002092//SPT23 PROTEIN. //0.12:208:26//SACCH 10 AROMYCES CEREVISIAE (BAKER'S YEAST). //P35210

F-HEMBA1002100

F-HEMBA1002102//ANKYRIN. //1.4e-12:106:35//MUS MUSCULUS (MOUSE). //Q02357

F-HEMBA1002113//EARLY NODULIN 20 PRECURSOR (N-20). //0.073:155:32//MEDICAGO TRUNCATULA (BARREL MEDI C). //P93329

F-HEMBA1002119//MALE SPECIFIC SPERM PROTEIN MST84D B. //0.85:22:36//DROSOPHILA MELANOGASTER (FRUIT FL Y). //Q01643

F-HEMBA1002125//GAG POLYPROTEIN [CONTAINS: CORE PR OTEINS P15, P12, P30]. //0.35:111:33//FELINE SARCOM A VIRUS (STRAIN SNYDER-THEILEN). //P03338

F-HEMBA1002139//HYPOTHETICAL 12.4 KD PROTEIN IN SE C17-QCR1 INTERGENIC REGION. //0.88:72:25//SACCHAROM YCES CEREVISIAE (BAKER'S YEAST). //P38192

F-HEMBA1002144

F-HEMBA1002150//THROMBOMODULIN (FETOMODULIN) (TM) (FRAGMENT). //4.8e-10:65:46//BOS TAURUS (BOVINE). // P06579

F-HEMBA1002151//A-AGGLUTININ ATTACHMENT SUBUNIT PR ECURSOR. //0.24:146:28//SACCHAROMYCES CEREVISIAE (B AKER'S YEAST). //P32323

F-HEMBA1002153//NADH-UBIQUINONE OXIDOREDUCTASE CHA IN 4L (EC 1.6.5.3). //0.93:58:25//APIS MELLIFERA (H ONEYBEE). //P34859

F-HEMBA1002160//!!!! ALU SUBFAMILY SP WARNING ENTR Y !!!!!//5.1e-21:94:65//HOMO SAPIENS (HUMAN). //P39193

F-HEMBA1002161//MYOSIN HEAVY CHAIN, CARDIAC MUSCLE 40 BETA ISOFORM. //1.4e-51:180:56//SUS SCROFA (PIG). // P79293

F-HEMBA1002162//!!!! ALU SUBFAMILY SP WARNING ENTR Y !!!!!//4.1e-40:102:75//HOMO SAPIENS (HUMAN). //P39193

F-HEMBA1002166//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.8e-13:133:45//HOMO SAPIENS (HUMAN). //P39188

F-HEMBA1002177//ZINC FINGER PROTEIN 142 (KIAA0236) (HA4654). //0.0014:153:26//HOMO SAPIENS (HUMAN). // 50

P52746

F-HEMBA1002185

F-HEMBA1002189//!!!! ALU SUBFAMILY SQ WARNING ENTR Y !!!!!//0.86:46:45//HOMO SAPIENS (HUMAN). //P39194

F-HEMBA1002191//MALE SPECIFIC SPERM PROTEIN MST84D C. //0.037:14:57//DROSOPHILA MELANOGASTER (FRUIT FL Y). //Q01644

F-HEMBA1002199

F-HEMBA1002204

F-HEMBA1002212//DUAL SPECIFICITY MITOGEN-ACTIVATED PROTEIN KINASE KINASEDSOR1 (EC 2.7.1.-) (DOWNSTRE AM OF RAF) (MAPKK). //3.2e-13:201:30//DROSOPHILA ME LANOGASTER (FRUIT FLY). //Q24324

F-HEMBA1002215//TESTIN 2 (TES2) [CONTAINS: TESTIN 1 (TES1)]. //1.1e-62:147:84//MUS MUSCULUS (MOUSE). // P47226

F-HEMBA1002226//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.6e-26:168:44//HOMO SAPIENS (HUMAN). //P39188

20 F-HEMBA1002229//!!!! ALU SUBFAMILY SB1 WARNING ENT RY !!!!!//6.8e-18:68:72//HOMO SAPIENS (HUMAN). //P39190

F-HEMBA1002237//EAMZP30-47 PROTEIN (FRAGMENT). //0.96:21:61//EIMERIA ACERVULINA. //P21959

F-HEMBA1002241//METALLOTHIONEIN (MT). //0.95:25:48//PARACENTROTUS LIVIDUS(COMMON SEA URCHIN). //P80367

F-HEMBA1002253//METALLOTHIONEIN-II (MT-II). //0.97:27:48//MESOCRICETUS AURATUS (GOLDEN HAMSTER). //P17808

30 F-HEMBA1002257

F-HEMBA1002265//MALE SPECIFIC SPERM PROTEIN MST84D C. //0.95:24:50//DROSOPHILA MELANOGASTER (FRUIT FL Y). //Q01644

F-HEMBA1002267//NEURONAL PROTEIN 3.1 (P311 PROTEI N). //0.94:33:33//GALLUSGALLUS (CHICKEN). //Q90667

F-HEMBA1002270

F-HEMBA1002321//HYPOTHETICAL IMMUNITY REGION PROTE IN 14. //0.99:22:40//BACTERIOPHAGE PHI-105. //P10437

F-HEMBA1002328

F-HEMBA1002337

F-HEMBA1002341//P53-BINDING PROTEIN 53BP2 (FRAGMEN T). //3.7e-55:109:96//MUS MUSCULUS (MOUSE). //Q62415

F-HEMBA1002348//PROBABLE E5 PROTEIN. //0.43:30:50// HUMAN PAPILLOMAVIRUS TYPE 35. //P27226

F-HEMBA1002349

F-HEMBA1002363//CHROMOSOME ASSEMBLY PROTEIN XCAP-E. //5.7e-105:278:71//XENOPUS LAEVIS (AFRICAN CLAWE D FROG). //P50533

F-HEMBA1002381//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//3.3e-24:69:73//HOMO SAPIENS (HUMAN). //P39188

- 8
F-HEMBA1002389//EARLY NODULIN 20 PRECURSOR (N-20).
//0.16:110:31//MEDICAGO TRUNCATULA (BARREL MEDIC).
//P93329
F-HEMBA1002417//TIGHT JUNCTION PROTEIN ZO-1 (TIGHT
JUNCTION PROTEIN 1).//2.6e-51:187:56//MUS MUSCULU
S (MOUSE).//P39447
F-HEMBA1002419//PROLINE-RICH PEPTIDE P-B.//1.0:18:
61//HOMO SAPIENS (HUMAN).//P02814
F-HEMBA1002430//HYPOTHETICAL 12.3 KD PROTEIN IN CA 10
PI-NAP1 INTERGENIC REGION.//0.042:41:46//SACCHAROM
YCES CEREVISIAE (BAKER'S YEAST).//P36140
F-HEMBA1002439//CHLOROPLAST 50S RIBOSOMAL PROTEIN
L27 (FRAGMENT).//0.99:47:29//CALYPTROSPHAERA SPHAE
ROIDEA.//P41548
F-HEMBA1002458//OVARIAN GRANULOSA CELL 13.0 KD PRO
TEIN HGR74.//4.1e-24:109:55//HOMO SAPIENS (HUMAN).
//Q00994
F-HEMBA1002460
F-HEMBA1002462//SALIVARY PROLINE-RICH PROTEIN II-1 20
(FRAGMENT).//0.00025:80:30//HOMO SAPIENS (HUMAN).
//P81489
F-HEMBA1002469//PUTATIVE TUMOR SUPPRESSOR LUCA15./
/0.0012:110:33//HOMO SAPIENS (HUMAN).//P52756
F-HEMBA1002475//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T).//0.37:106:33//MUSMUSCULUS (MOUSE).//P05143
F-HEMBA1002477//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/3.3e-34:96:71//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBA1002486
F-HEMBA1002495//LIGHT-MEDIATED DEVELOPMENT PROTEIN
DET1.//2.9e-31:110:39//ARABIDOPSIS THALIANA (MOUS
E-EAR CRESS).//P48732
F-HEMBA1002498//SFT2 PROTEIN.//1.0:54:35//SACCHARO
MYCES CEREVISIAE (BAKER'S YEAST).//P38166
F-HEMBA1002503//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/3.0e-06:49:63//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1002508//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/1.6e-22:169:44//HOMO SAPIENS (HUMAN).//P39
195
F-HEMBA1002513//HYPOTHETICAL 89.8 KD PROTEIN F41H1
0.6 IN CHROMOSOME IV.//0.00017:79:35//CAENORHABDIT
IS ELEGANS.//Q20296
F-HEMBA1002515
F-HEMBA1002538//ATP SYNTHASE E CHAIN, MITOCHONDRIA
L (EC 3.6.1.34).//1.0:53:37//SACCHAROMYCES CEREVIS
IAE (BAKER'S YEAST).//P81449
F-HEMBA1002542//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/5.7e-32:96:75//HOMO SAPIENS (HUMAN).//P3918 50
- 8
F-HEMBA1002547//AGRIN PRECURSOR.//2.5e-100:218:80/
/RATTUS NORVEGICUS (RAT).//P25304
F-HEMBA1002552//HEP27 PROTEIN (PROTEIN D).//9.5e-1
2:29:82//HOMO SAPIENS (HUMAN).//Q13268
F-HEMBA1002555//COLLAGEN ALPHA 1(III) CHAIN.//2.4e
-15:207:36//BOS TAURUS (BOVINE).//P04258
F-HEMBA1002558//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/1.0:34:50//HOMO SAPIENS (HUMAN).//P39193
F-HEMBA1002561//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//1.5e-05:49:46//NYCTICEBUS COUCANG (SLOW LORI
S).//P08548
F-HEMBA1002569//SINGLE-STRANDED DNA-BINDING PROTEI
N P12.//0.97:60:33//BACTERIOPHAGE PRD1.//P17637
F-HEMBA1002583
F-HEMBA1002590//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/5.6e-15:54:55//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1002592//HISTIDINE-RICH PROTEIN.//0.99:39:2
8//PLASMODIUM FALCIPARUM (ISOLATE FCM17 / SENEGA
L).//P14586
F-HEMBA1002609//SSM4 PROTEIN.//1.9e-12:135:29//SAC
CHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40318
F-HEMBA1002621//CYTOCHROME B6-F COMPLEX 3.5 KD SUB
UNIT (CYTOCHROME B6-FCOMPLEX SUBUNIT 6).//1.0:20:5
5//ZEA MAYS (MAIZE).//P19445
F-HEMBA1002624//EXTENSIN PRECURSOR (CELL WALL HYDR
OXYPROLINE-RICH GLYCOPROTEIN).//0.0035:124:33//NIC
OTIANA TABACUM (COMMON TOBACCO).//P13983
30 F-HEMBA1002628
F-HEMBA1002629//IMMEDIATE-EARLY PROTEIN IE180.//0.
84:80:36//PSEUDORABIESVIRUS (STRAIN KAPLAN) (PRV).
//P33479
F-HEMBA1002645//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/1.8e-16:57:68//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBA1002651
F-HEMBA1002659//CUTICLE COLLAGEN 2.//0.0077:77:38/
/CAENORHABDITIS ELEGANS.//P17656
40 F-HEMBA1002661//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//2.3e-89:116:72//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1002666//BETA CRYSTALLIN A4.//0.18:58:44//G
ALLUS GALLUS (CHICKEN).//P49152
F-HEMBA1002678
F-HEMBA1002679//GLUTAMIC ACID-RICH PROTEIN PRECURS
OR.//5.7e-06:219:27//PLASMODIUM FALCIPARUM (ISOLAT
E FC27 / PAPUA NEW GUINEA).//P13816
F-HEMBA1002688//SPIROIN 2 (DRAGLINE SILK FIBROIN
2) (FRAGMENT).//1.1e-07:198:32//NEPHILA CLAVIPES
(ORB SPIDER).//P46804

2879

F-HEMBA1002696//COLLAGEN ALPHA 1(VII) CHAIN PRECURSOR (LONG-CHAIN COLLAGEN) (LC COLLAGEN).//0.16:15
8:33//HOMO SAPIENS (HUMAN).//Q02388
F-HEMBA1002703//HYPOTHETICAL BHLF1 PROTEIN.//0.78:
147:29//EPSTEIN-BARR VIRUS (STRAIN B95-8) (HUMAN H
ERPESVIRUS 4).//P03181
F-HEMBA1002712//11.2 KD PROTEIN (ORF 103).//0.029:
75:34//BACTERIOPHAGE PF1.//P25133
F-HEMBA1002716//50S RIBOSOMAL PROTEIN L28.//1.0:4
4:27//BACILLUS SUBTILIS.//P37807
F-HEMBA1002728//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/5.4e-18:56:75//HOMO SAPIENS (HUMAN).//P391
95
F-HEMBA1002730//HYPOTHETICAL PROTEIN MJ0316.//0.09
7:84:35//METHANOCOCCUSJANNASCHII.//Q57764
F-HEMBA1002742//APOLIPOPROTEIN C-III PRECURSOR (AP
O-CIII).//0.97:26:50//SUS SCROFA (PIG).//P27917
F-HEMBA1002746//CALPHOTIN.//0.35:65:35//DROSOPHILA
MELANOGASTER (FRUIT FLY).//Q02910
F-HEMBA1002748//PLATELET GLYCOPROTEIN IB BETA CHAI
N PRECURSOR (GP-IB BETA) (GPIIB).//1.0:74:32//MUS
MUSCULUS (MOUSE).//P56400
F-HEMBA1002750//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/7.0e-15:49:75//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1002768//HYPOTHETICAL 72.2 KD PROTEIN C12C
2.05C IN CHROMOSOME II.//0.00036:197:26//SCHIZOSAC
CHAROMYCES POMBE (FISSION YEAST).//Q09746
F-HEMBA1002770//UTEROGLOBIN PRECURSOR (BLASTOKINI
N).//0.23:88:27//ORYCTOLAGUS CUNICULUS (RABBIT).// 30
P02779
F-HEMBA1002777//HOMEBOX PROTEIN HOX-A4 (HOX-1.4)
(MH-3).//0.00018:67:43//MUS MUSCULUS (MOUSE).//P06
798
F-HEMBA1002779//HYPOTHETICAL 17.6 KD PROTEIN IN NP
R1-RPS3 INTERGENIC REGION.//0.70:30:53//SACCHAROMY
CES CEREVISIAE (BAKER'S YEAST).//P53880
F-HEMBA1002780//OLFACTORY RECEPTOR 3 (K10) (FRAGME
NT).//1.0:31:45//MUS MUSCULUS (MOUSE).//Q60879
F-HEMBA1002794//HMG-Y RELATED PROTEIN B (SB16B PRO
TEIN) (FRAGMENT).//0.0044:66:37//GLYCINE MAX (SOYB
EAN).//Q10370
F-HEMBA1002801
F-HEMBA1002810//HYPOTHETICAL 25.9 KD PROTEIN AH6.3
IN CHROMOSOME II.//0.0033:116:31//CAENORHABDITIS
ELEGANS.//Q09202
F-HEMBA1002816//HYPOTHETICAL 47.1 KD PROTEIN C9G1.
13C IN CHROMOSOME I.//1.0e-17:68:48//SCHIZOSACCHAR
OMYCES POMBE (FISSION YEAST).//014308
F-HEMBA1002818//FIBULIN-2 PRECURSOR.//2.1e-27:92:4 50

2880

4//MUS MUSCULUS (MOUSE).//P37889
F-HEMBA1002826//DNA-BINDING PROTEIN 65 (PROTEIN GP
65).//0.28:46:34//BACTERIOPHAGE T4.//P16012
F-HEMBA1002833
F-HEMBA1002850//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//1.0:27:37//METRIDIDIUM SENILE (BROWN SEA
ANEMONE) (FRILLED SEA ANEMONE).//047493
F-HEMBA1002863//PHOTOSYSTEM I REACTION CENTRE SUBU
NIT IV (PHOTOSYSTEM 18.1 KD PROTEIN) (P30 PROTEIN)
10 (PSI-E).//0.84:37:43//SYNECHOCYSTIS SP. (STRAIN P
CC 6803).//P12975
F-HEMBA1002876//OCTAPEPTIDE-REPEAT PROTEIN T2.//0.
74:58:34//MUS MUSCULUS(MOUSE).//Q06666
F-HEMBA1002886
F-HEMBA1002896//HOMEBOX PROTEIN HOX-B3 (HOX-2G)
(HOX-2.7).//4.7e-05:84:35//HOMO SAPIENS (HUMAN).//
P14651
F-HEMBA1002921//TAT PROTEIN (TRANSACTIVATING REGUL
ATORY PROTEIN).//0.21:42:42//HUMAN IMMUNODEFICIENC
Y VIRUS TYPE 1 (RF/HAT ISOLATE) (HIV-1).//P05908
F-HEMBA1002924//EC PROTEIN HOMOLOG 2 (FRAGMENT).//
0.85:75:22//ARABIDOPSIS THALIANA (MOUSE-EAR CRES
S).//Q42377
F-HEMBA1002934//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/3.4e-31:92:72//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1002935//CASTRULA ZINC FINGER PROTEIN XLCCF
58.1 (FRAGMENT).//7.7e-06:187:29//XENOPUS LAEVIS
(AFRICAN CLAWED FROG).//P18730
F-HEMBA1002937//SUPPRESSOR PROTEIN SRP40.//0.0003
1:150:24//SACCHAROMYCESCEREVISIAE (BAKER'S YEAST).
//P32583
F-HEMBA1002939//ANKYRIN R (ANKYRINS 2.1 AND 2.2)
(ERYTHROCYTE ANKYRIN).//5.2e-25:225:33//HOMO SAPIE
NS (HUMAN).//P16157
F-HEMBA1002944
F-HEMBA1002951//TRICHOHYALIN.//0.0011:220:24//HOMO
SAPIENS (HUMAN).//Q07283
F-HEMBA1002954//PROBABLE E8 PROTEIN.//0.98:49:32//
BOVINE PAPILLOMAVIRUSTYPE 4.//P08352
F-HEMBA1002968//ACCESSORY GLAND PEPTIDE PRECURSOR
(PARAGONIAL PEPTIDE B).//0.93:41:34//DROSOPHILA SE
CHELLIA (FRUIT FLY).//018417
F-HEMBA1002970//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/0.00010:35:62//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBA1002971//INSULIN.//1.0:31:35//HYDROLAGUS CO
LLIEI (SPOTTED RATFISH) (PACIFIC RATFISH), AND CHI
MAERA MONSTROSA (RABBIT FISH).//P09536
F-HEMBA1002973//CAMP-DEPENDENT 3',5'-CYCLIC PHOSPH

2881

ODIESTERASE 4B (EC 3.1.4.17) (DPDE4). //3.0e-29:63:
100//RATTUS NORVEGICUS (RAT). //P14646
F-HEMBA1002997//HYPOTHETICAL 106.5 KD PROTEIN IN C
TT1-PRP31 INTERGENIC REGION. //1.0e-08:211:28//SACC
HAROMYCES CEREVISIAE (BAKER'S YEAST). //P53253
F-HEMBA1002999//SUPPRESSOR PROTEIN SRP40. //0.026:1
75:23//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32583
F-HEMBA1003021//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//1.3e-36:102:70//HOMO SAPIENS (HUMAN). //P39 10
194
F-HEMBA1003033//HYPOTHETICAL 23.1 KD PROTEIN CY27
7.20C. //0.029:75:29//MYCOBACTERIUM TUBERCULOSIS. //P71779
F-HEMBA1003034//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!//6.3e-23:144:46//HOMO SAPIENS (HUMAN). //P39
192
F-HEMBA1003035//HYPOTHETICAL 13.3 KD PROTEIN IN AR
OD-COMER INTERGENIC REGION. //0.99:55:30//BACILLUS
SUBTILIS. //P54457
F-HEMBA1003037//DNA-BINDING PROTEIN INHIBITOR ID-
4. //0.17:42:40//HOMO SAPIENS (HUMAN). //P47928
F-HEMBA1003041//DNA-BINDING P52/P100 COMPLEX, 100
KD SUBUNIT (FRAGMENTS). //0.69:28:46//HOMO SAPIENS
(HUMAN). //P30808
F-HEMBA1003046//MITOCHONDRIAL PROCESSING PROTEASE
BETA SUBUNIT PRECURSOR (EC 3.4.24.64) (BETA-MPP) (P
-52). //7.9e-124:253:96//HOMO SAPIENS (HUMAN). //075
439
F-HEMBA1003064//NADH-UBIQUINONE OXIDOREDUCTASE CHA
IN 5 (EC 1.6.5.3). //0.25:263:22//TRYPANOSOMA BRUCE
I BRUCEI. //P04540
F-HEMBA1003067//HYPOTHETICAL 57.5 KD PROTEIN IN VM
A7-RPS25A INTERGENIC REGION. //4.1e-05:189:30//SACC
HAROMYCES CEREVISIAE (BAKER'S YEAST). //P53214
F-HEMBA1003071//CUTICLE COLLAGEN 40. //6.0e-07:126:
38//CAENORHABDITIS ELEGANS. //P34804
F-HEMBA1003077//FIBROMODULIN PRECURSOR (FM) (COLLA
GEN-BINDING 59 KD PROTEIN). //2.4e-12:139:34//HOMO
SAPIENS (HUMAN). //Q06828
F-HEMBA1003078//RETROVIRUS-RELATED POL POLYPROTEIN
[CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); E
NDONUCLEASE]. //7.2e-05:60:40//MUS MUSCULUS (MOUS
E). //P11369
F-HEMBA1003079//PROTEIN Q300. //0.0012:16:87//MUS M
USCULUS (MOUSE). //Q02722
F-HEMBA1003083//!!!! ALU SUBFAMILY SB WARNING ENTR
Y !!!!!//3.3e-32:95:75//HOMO SAPIENS (HUMAN). //P391
89
F-HEMBA1003086

2882

F-HEMBA1003096//PROTAMINE IA (IRIDINE IA). //0.36:2
0:40//SALMO IRIDEUS (RAINBOW TROUT). //P02328
F-HEMBA1003098//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//4.4e-09:43:72//HOMO SAPIENS (HUMAN). //P3918
8
F-HEMBA1003117//PUTATIVE CUTICLE COLLAGEN C09G5.5.
//1.0:88:38//CAENORHABDITIS ELEGANS. //Q09456
F-HEMBA1003129//NADH-UBIQUINONE OXIDOREDUCTASE CHA
IN 4L (EC 1.6.5.3). //0.61:63:25//APIS MELLIFERA (H
ONEYBEE). //P34859
F-HEMBA1003133//COLLAGEN ALPHA 2(VIII) CHAIN (ENDO
THELIAL COLLAGEN) (FRAGMENT). //0.48:79:37//HOMO SA
PIENS (HUMAN). //P25067
F-HEMBA1003136//MANNOSYL-1-PHOSPHATE GUANYLTRANSFER
ASE (EC 2.7.7.13) (ATP-MANNOSYL-1-PHOSPHATE GUANYL
YLTRANSFERASE) (NDP-HEXOSE PYROPHOSPHORYLASE). //3.
6e-25:190:34//SACCHAROMYCES CEREVISIAE (BAKER'S YE
AST). //P41940
F-HEMBA1003142
20 F-HEMBA1003148//HYPOTHETICAL 56.4 KD PROTEIN IN RP
L30-CWH41 INTERGENIC REGION PRECURSOR. //0.068:171:
23//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53
189
F-HEMBA1003166//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!//1.8e-13:54:66//HOMO SAPIENS (HUMAN). //P391
92
F-HEMBA1003175//HYPOTHETICAL 57.5 KD PROTEIN IN VM
A7-RPS25A INTERGENIC REGION. //0.015:147:31//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST). //P53214
30 F-HEMBA1003179//PROBABLE TRNA (5-METHYLAMINOMETHYL
-2-THIOURIDYLATE)-METHYLTRANSFERASE (EC 2.1.1.61).
//2.6e-51:164:47//BACILLUS SUBTILIS. //035020
F-HEMBA1003197
F-HEMBA1003199//HOMEODOMAIN PROTEIN HOX-A4 (HOX-1D)
(HOX-1.4). //0.00049:83:38//HOMO SAPIENS (HUMAN). //Q00056
F-HEMBA1003202//SPERM PROTAMINE P1. //0.98:53:28//P
LANIGALE GILESI (FLAT-SKULLED MARSUPIAL MOUSE). //O
18747
40 F-HEMBA1003204//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//5.2e-22:42:80//HOMO SAPIENS (HUMAN). //P391
94
F-HEMBA1003212//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!//1.6e-18:74:71//HOMO SAPIENS (HUMAN). //P391
93
F-HEMBA1003220//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//3.3e-18:56:78//HOMO SAPIENS (HUMAN). //P391
94
F-HEMBA1003222//HYPOTHETICAL 37.5 KD PROTEIN IN GN
TR-HTPG INTERGENIC REGION. //0.0018:159:27//BACILLUS

2883

2884

S SUBTILIS.//P46327

F-HEMBA1003229//DIHYDRODIPICOLINATE SYNTHASE 1 PRECURSOR (EC 4.2.1.52) (DHDPS).//1.0:85:28//TRITICUM AESTIVUM (WHEAT).//P24846

F-HEMBA1003235//TROPOMYOSIN.//8.3e-07:109:33//SCHIZOSACCHAROMYCES POMBE(FISSION YEAST).//Q02088

F-HEMBA1003250

F-HEMBA1003257//HYPOTHETICAL PROLINE-RICH PROTEIN (FRAGMENT).//1.5e-07:27:74//OWENIA FUSIFORMIS.//P21260

F-HEMBA1003273

F-HEMBA1003276

F-HEMBA1003278

F-HEMBA1003281//HOMEBOX PROTEIN HOX-A4 (CHOX-1.4).//0.0053:116:36//GALLUS GALLUS (CHICKEN).//P17277

F-HEMBA1003286//DNA-DIRECTED RNA POLYMERASE SUBUNIT N (EC 2.7.7.6).//0.96:37:35//SULFOLOBUS ACIDOCALDARIUS.//P39472

F-HEMBA1003291//5'-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC ALPHA-2 CHAIN(EC 2.7.1.-) (AMPK ALPHA-2 CHAIN) (FRAGMENT).//3.3e-15:68:39//SUS SCROFA (PIG).//Q28948

F-HEMBA1003296//PULMONARY SURFACTANT-ASSOCIATED PROTEIN B (SP-B) (6 KD PROTEIN) (PULMONARY SURFACTANT-ASSOCIATED PROTEOLIPID SPL(PHE)).//0.98:49:28//BOS TAURUS (BOVINE).//P15781

F-HEMBA1003304//MITOCHONDRIAL RIBOSOMAL PROTEIN S19.//0.99:36:30//PROTOHECA WICKERHAMII.//P46750

F-HEMBA1003309//HYPOTHETICAL 7.9 KD PROTEIN.//0.69:54:37//VACCINIA VIRUS(STRAIN WR), AND VACCINIA VIRUS (STRAIN COPENHAGEN).//P04306

F-HEMBA1003314//MIXED LINEAGE KINASE 2 (EC 2.7.1.-) (FRAGMENT).//2.3e-06:143:22//HOMO SAPIENS (HUMAN).//Q02779

F-HEMBA1003322//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//1.5e-30:53:77//HOMO SAPIENS (HUMAN).//P39194

F-HEMBA1003327

F-HEMBA1003328//TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN) (FRAGMENT).//0.53:21:42//HUMAN IMMUNODEFICIENCY VIRUS TYPE 1 (BH5 ISOLATE) (HIV-1).//P04612

F-HEMBA1003330//LONG NEUROTOXIN 3 (TOXIN VN2).//1.0:26:34//DENDROASPIS POLYLEPIS POLYLEPIS (BLACK MAMBIA).//P25667

F-HEMBA1003348//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//6.5e-09:56:66//HOMO SAPIENS (HUMAN).//P39194

F-HEMBA1003369//ANTER-SPECIFIC PROLINE-RICH PROTEIN

N APG PRECURSOR.//0.0042:97:36//ARABIDOPSIS THALIANA (MOUSE-EAR CRESS).//P40602

F-HEMBA1003370//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//7.0e-18:99:53//HOMO SAPIENS (HUMAN).//P39188

F-HEMBA1003373

F-HEMBA1003376//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//4.7e-16:60:75//HOMO SAPIENS (HUMAN).//P39189

10 F-HEMBA1003380//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//2.8e-10:50:68//HOMO SAPIENS (HUMAN).//P39188

F-HEMBA1003384

F-HEMBA1003395//PROBABLE E5 PROTEIN.//0.62:64:29//HUMAN PAPILLOMAVIRUS TYPE 16.//P06927

F-HEMBA1003402//HYPOTHETICAL 12.0 KD PROTEIN IN TUBULIN-INTERGENIC REGION PRECURSOR.//0.89:74:33//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q04521

F-HEMBA1003403//ACIDIC PROLINE-RICH PROTEIN PRECURSOR (CLONE PRP25) (FRAGMENT).//0.0010:69:33//RATTUS NORVEGICUS (RAT).//P10164

F-HEMBA1003408//WEB1 PROTEIN (PROTEIN TRANSPORT PROTEIN SEC31).//4.8e-06:93:25//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38968

F-HEMBA1003417//PROCOLLAGEN ALPHA 1(II) CHAIN PRECURSOR [CONTAINS: CHONDROCALCIN].//0.0021:140:34//MUS MUSCULUS (MOUSE).//P28481

F-HEMBA1003418//SPLICING FACTOR, ARGININE/SERINE-RICH 4 (PRE-MRNA SPLICING FACTOR SRP75).//1.7e-14:188:33//HOMO SAPIENS (HUMAN).//Q08170

F-HEMBA1003433//DNA REPAIR PROTEIN XRS2.//1.0:88:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P33301

F-HEMBA1003447//ACIDIC PROLINE-RICH PROTEIN PRECURSOR (CLONE PRP25) (FRAGMENT).//0.0061:69:33//RATTUS NORVEGICUS (RAT).//P10164

F-HEMBA1003461//SPIDROIN 1 (DRAGLINE SILK FIBROIN 1) (FRAGMENT).//2.3e-09:239:33//NEPHILA CLAVIPES (ORB SPIDER).//P19837

F-HEMBA1003463//METALLOTHIONEIN-A (MTA) (FRAGMENT).//1.0:40:35//SPHAERECHINUS GRANULARIS (PURPLE SEA URCHIN).//Q26497

F-HEMBA1003480//FUSARIC ACID RESISTANCE PROTEIN FUSB.//0.0043:96:32//BURKHOLDERIA CEPACIA (PSEUDOMONAS CEPACIA).//P24127

F-HEMBA1003528//36.4 KD PROLINE-RICH PROTEIN.//6.4e-15:167:33//LYCOPERSICON ESCULENTUM (TOMATO).//Q0451

F-HEMBA1003531//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//1.2e-18:56:78//HOMO SAPIENS (HUMAN).//P391

- 89
F-HEMBA1003538//COMPLEMENT C1R COMPONENT PRECURSOR (EC 3.4.21.41).//2.5e-28:136:47//HOMO SAPIENS (HUMAN).//P00736
F-HEMBA1003545//INSULIN GENE ENHANCER PROTEIN ISL-2 (ISLET-2).//9.2e-105:217:85//RATTUS NORVEGICUS (RAT).//P50480
F-HEMBA1003548
F-HEMBA1003555//HYPOTHETICAL 31.9 KD PROTEIN IN BE T1-PAN1 INTERGENIC REGION.//8.7e-57:180:55//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40558
F-HEMBA1003556//HYPOTHETICAL 19.2 KD PROTEIN IN CO X-REP INTERGENIC REGION (ORF5) (ORF21).//0.53:97:25//BACTERIOPHAGE HP1.//P51706
F-HEMBA1003560//GUANINE NUCLEOTIDE-BINDING PROTEIN G(1)/G(S)/G(O) GAMMA-2 SUBUNIT (G GAMMA-I).//1.8e-32:71:100//BOS TAURUS (BOVINE).//P16874
F-HEMBA1003568//ZINC-FINGER PROTEIN RFP (RET FINGER PROTEIN).//4.1e-19:126:31//HOMO SAPIENS (HUMAN).//P14373
F-HEMBA1003569//METASTASIS-ASSOCIATED PROTEIN MTA1.//3.9e-83:143:74//HOMO SAPIENS (HUMAN).//Q13330
F-HEMBA1003571//HYPOTHETICAL 8.7 KD PROTEIN (READING FRAME D).//1.0:64:25//STAPHYLOCOCCUS AUREUS.//P03860
F-HEMBA1003579//CYTOTOXIN 1 (CYTOTOXIN V-II-1) (TOXIN V(II)1).//1.0:41:29//NAJA MELANOLEUCA (FOREST COBRA) (BLACK-LIPPED COBRA).//P01448
F-HEMBA1003581//TALIN.//3.7e-36:52:98//MUS MUSCULUS (MOUSE).//P26039
F-HEMBA1003591//CHLOROPLAST 28 KD RIBONUCLEOPROTEIN PRECURSOR (28RNP).//1.6e-05:91:31//NICOTIANA SYLVESTRIS (WOOD TOBACCO).//P19682
F-HEMBA1003595//HYPOTHETICAL 12.0 KD PROTEIN IN DS T1-HEM2 INTERGENIC REGION.//1.0:55:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53182
F-HEMBA1003597
F-HEMBA1003598//T-CELL RECEPTOR BETA CHAIN PRECURSOR (ANA 11).//4.9e-10:85:41//ORYZOLAGUS CUNICULUS (RABBIT).//P06333
F-HEMBA1003615//PUTATIVE MINOR COAT PROTEIN (ORF43).//0.086:10:70//BACTERIOPHAGE PHI-LF.//Q07482
F-HEMBA1003617//HYPOTHETICAL 36.8 KD PROTEIN C26A3.16 IN CHROMOSOME I.//4.4e-13:58:48//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q10169
F-HEMBA1003621//LONG NEUROTOXIN 1 (NEUROTOXIN A).//0.096:40:37//OPHIOPHAGUS HANNAH (KING COBRA) (NAJA HANNAH).//P01387
F-HEMBA1003622
F-HEMBA1003630
F-HEMBA1003637//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//2.4e-13:47:74//HOMO SAPIENS (HUMAN).//P39188
8
F-HEMBA1003640//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//0.87:25:64//HOMO SAPIENS (HUMAN).//P39193
F-HEMBA1003645//HYPOTHETICAL 40.4 KD TRP-ASP REPEATS CONTAINING PROTEIN C14B1.4 IN CHROMOSOME III.//1.8e-10:157:26//CAENORHABDITIS ELEGANS.//Q17963
F-HEMBA1003646//SERINE-ARGININE PROTEIN 55 (SRP55) (ENHANCER OF DEFORMED) (52-KD BRACKETING PROTEIN) (B52 PROTEIN).//4.9e-05:207:27//DROSOPHILA MELANOGASTER (FRUIT FLY).//P26686
F-HEMBA1003656
F-HEMBA1003662//PROLINE-RICH PEPTIDE P-B.//0.57:17:52//HOMO SAPIENS (HUMAN).//P02814
F-HEMBA1003667//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//6.0e-16:43:72//HOMO SAPIENS (HUMAN).//P39194
F-HEMBA1003679
20 F-HEMBA1003680//PUTATIVE AMINOPEPTIDASE ZK353.6 IN CHROMOSOME III (EC 3.4.11.-).//3.9e-08:137:27//CAENORHABDITIS ELEGANS.//P34629
F-HEMBA1003684//ZINC FINGER PROTEIN 151 (POLYOMAVIRUS LATE INITIATOR PROMOTER BINDING PROTEIN) (LP-1) (ZINC FINGER PROTEIN Z13).//2.1e-20:127:40//MUS MUSCULUS (MOUSE).//Q60821
F-HEMBA1003690//HYPOTHETICAL PROTEIN KIAA0288 (HA6116).//3.0e-85:201:78//HOMO SAPIENS (HUMAN).//P56524
30 F-HEMBA1003692//CELL DIVISION CONTROL PROTEIN 1.//0.13:69:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40986
F-HEMBA1003711//CARCINOEMBRYONIC ANTIGEN PRECURSOR (CEA) (MECONIUM ANTIGEN 100) (CD66E ANTIGEN).//0.021:153:26//HOMO SAPIENS (HUMAN).//P06731
F-HEMBA1003714//ABAECIN.//0.99:34:32//BOMBUS PASCORUM.//P81463
F-HEMBA1003715
F-HEMBA1003720//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//5.4e-34:155:56//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1003725//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//1.3e-27:181:41//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1003729//PTB-ASSOCIATED SPLICING FACTOR (PSF).//0.0037:103:33//HOMO SAPIENS (HUMAN).//P23246
F-HEMBA1003733//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//5.0e-54:210:58//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1003742//MALE SPECIFIC SPERM PROTEIN MST84DB.//0.066:72:33//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01643
50 F-HEMBA1003758

2887

2888

- F-HEMBA1003760//HYPOXIA-INDUCIBLE FACTOR 1 ALPHA (HIF-1 ALPHA) (ARNT INTERACTING PROTEIN).//1.5e-5
1:220:52//MUS MUSCULUS (MOUSE).//Q61221
F-HEMBA1003773
F-HEMBA1003783
F-HEMBA1003784
F-HEMBA1003799//SHORT NEUROTOXIN 1 (TOXIN AA C).//
0.95:27:37//ACANTHOPHIS ANTARCTICUS (COMMON DEATH
ADDER).//P01434
F-HEMBA1003803//GAG POLYPROTEIN [CONTAINS: CORE PR
OTEINS P15, P12, P30].//0.46:96:34//FELINE SARCOMA
VIRUS (STRAIN SNYDER-THEILEN).//P03338
F-HEMBA1003804//DNA-BINDING P52/P100 COMPLEX, 100
KD SUBUNIT (FRAGMENTS).//0.019:30:50//HOMO SAPIENS
(HUMAN).//P30808
F-HEMBA1003805//HYPOTHETICAL 75.0 KD PROTEIN B028
0.11 IN CHROMOSOME III.//1.8e-20:109:47//CAENORHAB
DITIS ELEGANS.//P42083
F-HEMBA1003807
F-HEMBA1003827//HYPOTHETICAL PROLINE-RICH PROTEIN 20
(FRAGMENT).//2.1e-09:23:78//OWENIA FUSIFORMIS.//P2
1260
F-HEMBA1003836//MOB1 PROTEIN (MPS1 BINDER 1).//2.0
e-31:134:52//SACCHAROMYCES CEREVISIAE (BAKER'S YEA
ST).//P40484
F-HEMBA1003838//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!/3.9e-22:39:76//HOMO SAPIENS (HUMAN).//P391
92
F-HEMBA1003856
F-HEMBA1003864//HYPOTHETICAL 39.4 KD PROTEIN IN ME 30
T1-SIS2 INTERGENIC REGION.//1.5e-15:194:30//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P36151
F-HEMBA1003866//PROTEIN A39.//0.0027:72:33//VACCIN
IA VIRUS (STRAIN COPENHAGEN).//P21062
F-HEMBA1003879//80 KD NUCLEAR CAP BINDING PROTEIN
(NCBP 80 KD SUBUNIT) (CBP80).//2.9e-16:22:100//HOM
O SAPIENS (HUMAN).//Q09161
F-HEMBA1003880//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//0.99:39:38//FELIS SILVESTRIUS CATUS (CA
T).//P48896
F-HEMBA1003885//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/3.5e-28:47:76//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBA1003893//HYPOTHETICAL 27.8 KD PROTEIN IN VM
A7-RPS25A INTERGENIC REGION.//1.7e-57:215:51//SACC
HAROMYCES CEREVISIAE (BAKER'S YEAST).//P53215
F-HEMBA1003902
F-HEMBA1003908
F-HEMBA1003926//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/4.3e-10:60:63//HOMO SAPIENS (HUMAN).//P3918 50
- 8
F-HEMBA1003937//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/8.1e-29:68:64//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBA1003939//PROTEIN Q300.//0.0025:24:62//MUS M
USCULUS (MOUSE).//Q02722
F-HEMBA1003942//EXCITATORY INSECT TOXIN BJXTR-IT P
RECURSOR (BJ-XTRIT).//0.084:67:31//BUTHOTUS JUDAIC
US (SCORPION) (HOTTENTOTTA JUDAICA).//P56637
F-HEMBA1003950//HYPOTHETICAL 8.1 KD PROTEIN IN SPE
A-METK INTERGENIC REGION (071).//0.95:26:34//ESCHE
RICHIA COLI.//P46878
F-HEMBA1003953//ZINC FINGER PROTEIN MFG-1 (ZINC FI
NGER PROTEIN 58) (FRAGMENT).//2.5e-17:89:46//MUS M
USCULUS (MOUSE).//P16372
F-HEMBA1003958//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//4.2e-23:43:76//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1003959
F-HEMBA1003976//HYPOTHETICAL PROTEIN KIAA0076 (HAO
936).//0.99:88:28//HOMO SAPIENS (HUMAN).//Q14999
F-HEMBA1003978//SMALL PROLINE RICH PROTEIN II (SPR
-II) (CLONE 930).//0.98:19:57//HOMO SAPIENS (HUMA
N).//P22531
F-HEMBA1003985//LYSYL-TRNA SYNTHETASE (EC 6.1.1.6)
(LYSINE--TRNA LIGASE) (LYSRS) (FRAGMENT).//1.0:40:
32//MYCOBACTERIUM LEPRAE.//P46861
F-HEMBA1003987//HYPOTHETICAL PROTEIN UL66.//0.27:6
5:33//HUMAN CYTOMEGALOVIRUS (STRAIN AD169).//P1682
2
F-HEMBA1003989//MALE SPECIFIC SPERM PROTEIN MST84D
B.//5.2e-05:64:40//DROSOPHILA MELANOGASTER (FRUIT
FLY).//Q01643
F-HEMBA1004000//PROTEIN Q300.//0.00042:17:82//MUS
MUSCULUS (MOUSE).//Q02722
F-HEMBA1004011//ALPHA-TYPE CALCITONIN GENE-RELATED
PEPTIDE PRECURSOR (CGRP-I).//0.47:106:32//HOMO SA
PIENS (HUMAN).//P06881
F-HEMBA1004012//ATP SYNTHASE PROTEIN 9, MITOCHONDR
IAL (EC 3.6.1.34) (LIPID-BINDING PROTEIN).//0.96:3
6:33//PARAMECIUM TETRAURELIA.//P16001
F-HEMBA1004015//HYPOTHETICAL 29.3 KD PROTEIN B028
0.6 IN CHROMOSOME III.//0.00018:90:34//CAENORHABDI
TIS ELEGANS.//P41997
F-HEMBA1004024//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/5.1e-34:75:80//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBA1004038
F-HEMBA1004042
F-HEMBA1004045//40S RIBOSOMAL PROTEIN S27A.//1.0:2
0:55//ASPARAGUS OFFICINALIS (GARDEN ASPARAGUS).//P

31753

F-HEMBA1004048//PROLINE-RICH PROTEIN MP-3 (FRAGMENT). //1.3e-06:158:35//MUS MUSCULUS (MOUSE). //P05143
F-HEMBA1004049//32 KD HEAT SHOCK PROTEIN (4-1 PROTEIN). //0.098:106:32//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD). //P54658

F-HEMBA1004055//HYPOTHETICAL PROTEIN HI0258/259. //0.87:133:23//HAEMOPHILUS INFLUENZAE. //P43974

F-HEMBA1004056//!!!! ALU SUBFAMILY SB2 WARNING ENTRY !!!!!//3.3e-25:39:64//HOMO SAPIENS (HUMAN). //P39191

F-HEMBA1004074//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//9.9e-08:35:68//HOMO SAPIENS (HUMAN). //P39188

F-HEMBA1004086F-HEMBA1004097//IMMEDIATE-EARLY PROTEIN IE4 (IE68) (FRAGMENT). //0.71:95:35//HERPES SIMPLEX VIRUS (TYPE 2). //P14379

F-HEMBA1004111//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//4.7e-26:84:64//HOMO SAPIENS (HUMAN). //P39188

F-HEMBA1004131//SEPTIN 2 HOMOLOG (FRAGMENT). //2.8e-34:108:63//HOMO SAPIENS (HUMAN). //Q14141

F-HEMBA1004132//HYPOTHETICAL PROTEIN HI1736. //1.0:44:34//HAEMOPHILUS INFLUENZAE. //P44300

F-HEMBA1004133//HYPOTHETICAL 8.5 KD PROTEIN CY274.40C. //0.89:21:57//MYCOBACTERIUM TUBERCULOSIS. //Q10826

F-HEMBA1004138//EARLY NODULIN 75 (N-75) (NGM-75) (FRAGMENT). //0.016:39:41//MEDICAGO SATIVA (ALFALFA). //P11728

F-HEMBA1004143//CYTOCHROME C OXIDASE POLYPEPTIDE V III PRECURSOR (EC 1.9.3.1). //0.93:34:29//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P04039

F-HEMBA1004146//BASIC PROLINE-RICH PEPTIDE P-E (IB 50

-9). //0.63:52:36//HOMO SAPIENS (HUMAN). //P02811

F-HEMBA1004150//METALLOTHIONEIN-II (MT-II). //1.0:20:45//MUS MUSCULUS (MOUSE). //P02798

F-HEMBA1004164//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//3.0e-13:57:71//HOMO SAPIENS (HUMAN). //P39195

F-HEMBA1004168//V-TYPE SODIUM ATP SYNTHASE SUBUNIT F (EC 3.6.1.34) (NA(+)-TRANSLOCATING ATPASE SUBUNIT F). //0.00035:90:34//ENTEROCOCCUS HIRAE. //P43437

F-HEMBA1004199//HYPOTHETICAL HELICASE K12H4.8 IN CHROMOSOME III. //5.1e-14:115:31//CAENORHABDITIS ELEGANS. //P34529

F-HEMBA1004200

F-HEMBA1004202//YPT1-RELATED PROTEIN 1. //2.5e-24:96:52//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //P11620

F-HEMBA1004203//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//2.2e-09:48:64//HOMO SAPIENS (HUMAN). //P39193

F-HEMBA1004207//HYPOTHETICAL 8.7 KD PROTEIN IN RPL22-RPL23 INTERGENIC REGION (ORF70). //0.98:51:33//ASTASIA LONGA (EUGLENOPHYCEAN ALGA). //P34779

F-HEMBA1004225//METALLOTHIONEIN-II. //1.0:30:33//CANDIDA GLABRATA (YEAST) (TORULOPSIS GLABRATA). //P15114

F-HEMBA1004227//PUTATIVE PROTEIN PHOSPHATASE 2C (EC 3.1.3.16) (PP2C) (KIAA0015). //5.9e-06:109:33//HOMO SAPIENS (HUMAN). //P49593

2891

2892

F-HEMBA1004238//VERY HYPOTHETICAL XYLU PROTEIN. //0.98:39:38//ESCHERICHIA COLI. //P05056

F-HEMBA1004241//SOX-13 PROTEIN (FRAGMENT). //0.66:36:38//MUS MUSCULUS (MOUSE). //Q04891

F-HEMBA1004246

F-HEMBA1004248//INSULIN-INDUCED GROWTH RESPONSE PROTEIN CL-6 (IMMEDIATE-EARLY PROTEIN CL-6). //1.0e-43:98:84//RATTUS NORVEGICUS (RAT). //Q08755

F-HEMBA1004264//SPIDROIN 2 (DRAGLINE SILK FIBROIN 2) (FRAGMENT). //0.014:160:28//NEPHILA CLAVIPES (ORBIT SPIDER). //P46804

F-HEMBA1004267//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//1.8e-52:56:83//HOMO SAPIENS (HUMAN). //P39189

F-HEMBA1004272

F-HEMBA1004274//HYPOTHETICAL 13.0 KD PROTEIN F59B. 2.10 IN CHROMOSOME III. //0.00084:33:54//CAENORHABDITIS ELEGANS. //P34485

F-HEMBA1004275//HYPOTHETICAL 56.5 KD PROTEIN IN CAJ1-HOM3 INTERGENIC REGION. //9.3e-06:125:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40034
F-HEMBA1004276//BETA-ADAPTIN 1 (PLASMA MEMBRANE ADAPTOR HA2/AP2 ADAPTIN BETA SUBUNIT) (CLATHRIN ASSEMBLY PROTEIN COMPLEX 2 BETA LARGE CHAIN) (AP105A). //3.7e-30:239:32//HOMO SAPIENS (HUMAN). //Q10567

F-HEMBA1004286//CUTICLE COLLAGEN 34. //0.0027:71:38//CAENORHABDITIS ELEGANS. //P34687

F-HEMBA1004289//PTR3 PROT 50

EIN (SSY3 PROTEIN). //1.0:76:28//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P43606

F-HEMBA1004295//DNA-BINDING P52/P100 COMPLEX, 100 KD SUBUNIT (FRAGMENTS). //0.075:58:39//HOMO SAPIENS (HUMAN). //P30808

10 F-HEMBA1004306//HYPOTHETICAL 29.3 KD PROTEIN (ORF92). //0.020:132:30//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPMNPV). //O10341

F-HEMBA1004312//EARLY PROTEIN I73R. //0.99:65:32//AFRICAN SWINE FEVER VIRUS (STRAIN BA71V) (ASFV). //P27946

20 F-HEMBA1004321//ZINC FINGER PROTEIN 90 (ZFP-90) (ZINC FINGER PROTEIN NK10). //4.3e-43:133:44//MUS MUSCULUS (MOUSE). //Q61967

F-HEMBA1004323

F-HEMBA1004327//SMALL PROLINE-RICH PROTEIN 2-1. //0.027:48:43//HOMO SAPIENS (HUMAN). //P35326

30 F-HEMBA1004330//HOMEBOX PROTEIN ENGRAILED-1 (HUN-1). //0.46:70:34//HOMO SAPIENS (HUMAN). //Q05925

F-HEMBA1004334//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //7.7e-05:83:34//HOMO SAPIENS (HUMAN). //P08547

F-HEMBA1004335//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//2.0e-24:41:80//HOMO SAPIENS (HUMAN). //P39195

F-HEMBA1004341//PROLINE-RICH PROTEIN MP-3 (FRAGMENT). //2.8e-06:148:35//MUS MUSCULUS (MOUSE). //P05143

F-HEMBA1004353//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//2.2e-29:57:80//HO

2893

MO SAPIENS (HUMAN). //P39195
 F-HEMBA1004354//CHL1 PROTEIN. //0.017:40:40//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P22516
 F-HEMBA1004356
 F-HEMBA1004366//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //0.00045:49:46//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1004372//VERY HYPOTHETICAL 20.6 KD PROTEIN C56F8.15 IN CHROMOSOME I. //1.0:125:28//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10263
 F-HEMBA1004389//HYPOTHETICAL 113.1 KD PROTEIN IN PRE5-FET4 INTERGENIC REGION. //0.76:170:25//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q04893
 F-HEMBA1004394
 F-HEMBA1004396//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //1.2e-10:72:51//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1004405
 F-HEMBA1004408//PEPTIDYL-PROLYL CIS-TRANS ISOMERASE 10 (EC 5.2.1.8) (PPIASE) (ROTAMASE) (CYCLOPHILIN-10). //2.7e-29:146:48//CAENORHABDITIS ELEGANS. //P52017
 F-HEMBA1004429//!!!! ALU SUBFAMILY SB2 WARNING ENTRY !!!!!//0.0019:47:59//HOMO SAPIENS (HUMAN). //P39191
 F-HEMBA1004433//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!//1.1e-20:47:68//HOMO SAPIENS (HUMAN). //P39192
 F-HEMBA1004460//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//6.2e-64:134:69//HOMO SAPIENS (HUMAN). //P39193
 F-HEMBA1004461//METALLOTHIONEIN-LIKE PROTEIN 1. //1.0:39:35//PISUM SATIVUM (GARDEN PEA). //P20830
 F-HEMBA1004479//HYPOXIA-INDUCIBLE FACTOR 1 ALPHA (HIF-1 ALPHA) (ARNT INTERACTING PROTEIN). //9.7e-43:101:48//MUS MUSCULUS (MOUSE). //Q61221
 F-HEMBA1004482//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34). //1.0:41:36//CANDIDA GLABRATA (YEAST) (TORULOPSIS GLABRATA). //P05040
 F-HEMBA1004499//TUBULIN BETA CHAIN. //0.00021:55:36//CAENORHABDITIS ELEGANS. //P52275
 F-HEMBA1004502
 F-HEMBA1004506//HYPOTHETICAL PROTEIN ORF-1137. //5.3e-11:119:35//MUS MUSCULUS (MOUSE). //P11260

2894

F-HEMBA1004507//SPLICEOSOME ASSOCIATED PROTEIN 62 (SAP 62) (SF3A66). //0.00072:90:37//HOMO SAPIENS (HUMAN). //Q15428
 F-HEMBA1004509//HYPOTHETICAL 52.2 KD PROTEIN IN MP R1-GCN20 INTERGENIC REGION. //6.3e-28:169:42//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P43589
 F-HEMBA1004534//ENDOTHELIAL ACTIN-BINDING PROTEIN (ABP-280) (NONMUSCLE FILAMIN) (FILAMIN 1). //1.3e-80:226:66//HOMO SAPIENS (HUMAN). //P21333
 10 F-HEMBA1004538//HYPOTHETICAL PROTEIN MJ0764. //0.96:28:35//METHANOCOCCUS JANNASCHII. //Q58174
 F-HEMBA1004542//METALLOTHIONEIN (MT). //0.78:36:41//GADUS MORHUA (ATLANTIC COD). //P51902
 F-HEMBA1004554
 F-HEMBA1004560//HYPOTHETICAL PROTEIN KIAA0281 (HA6725). //4.2e-15:56:69//HOMO SAPIENS (HUMAN). //Q92556
 F-HEMBA1004573//CIRCUMSPOROZOITE PROTEIN PRECURSOR (CS). //0.65:31:58//PLASMODIUM BERGHEI. //P06915
 20 F-HEMBA1004577//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//3.9e-08:35:80//HOMO SAPIENS (HUMAN). //P39195
 F-HEMBA1004586//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//6.6e-08:64:54//HOMO SAPIENS (HUMAN). //P39194
 F-HEMBA1004596//HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN C (HNRNP C) (HNRNP CORE PROTEIN C) (FRAGMENT). //0.00057:88:31//RATTUS NORVEGICUS (RAT). //P17132
 30 F-HEMBA1004604//COLLAGEN ALPHA 2(XI) CHAIN PRECURSOR (FRAGMENT). //0.045:37:45//MUS MUSCULUS (MOUSE). //Q64739
 F-HEMBA1004610//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//4.3e-11:73:54//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1004617
 F-HEMBA1004629
 F-HEMBA1004631//HYPOTHETICAL 7.8 KD PROTEIN IN WAP A-L1CT INTERGENIC REGION. //1.0:36:38//BACILLUS SUBTILIS. //P42303
 F-HEMBA1004632//PHOTOSYSTEM I REACTION CENTRE SUBUNIT X PRECURSOR (LIGHT-HARVESTING 8.0 KD POLYPEPTIDE). //0.86:48:35//SYNECHOCOCCUS ELONGATUS NAEGELI. //P20453
 F-HEMBA1004637//HYPOTHETICAL 83.6 KD PROTEIN R05D3.2 IN CHROMOSOME III. //1.7e-32:159:42//CAENORHABDITIS ELEGANS. //P34535
 F-HEMBA1004638//HYPOTHETICAL PROLINE-RICH PROTEIN (FRAGMENT). //2.8e-06:50:46//OWENIA FUSIFORMIS. //P21260
 50 1260

2895

F-HEMBA1004666//TOXIN S6C4.//1.0:36:30//DENDROASPIS JAMESONI KAIMOSAE (EASTERN JAMESON'S MAMBA).//P25682

F-HEMBA1004669//SPLICING FACTOR, ARGININE/SERINE-RICH 4 (PRE-MRNA SPLICING FACTOR SRP75).//1.6e-12:105:42//HOMO SAPIENS (HUMAN).//Q08170

F-HEMBA1004670//PROCOLLAGEN ALPHA 1(I) CHAIN PRECURSOR.//2.5e-06:62:45//HOMO SAPIENS (HUMAN).//P02452

F-HEMBA1004672//HYPOTHETICAL PROTEIN MJ0437.//0.95:37:29//METHANOCOCCUS JANNASCHII.//Q57879

F-HEMBA1004693//MYOSIN HEAVY CHAIN, NONMUSCLE TYPE B (CELLULAR MYOSIN HEAVY CHAIN, TYPE B) (NMHC-B).//0.00035:217:23//HOMO SAPIENS (HUMAN).//P35580
F-HEMBA1004697//IMMUNOGLOBULIN G BINDING PROTEIN H PRECURSOR (PROTEIN H).//0.058:118:30//STREPTOCOCCUS PYOGENES.//P50470

F-HEMBA1004705//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//6.8e-09:43:72//HOMO SAPIENS (HUMAN).//P39188

F-HEMBA1004709//!!!! ALU SUBFAMILY SB WARNING ENTRY
!!!!//8.8e-18:50:84//HOMO SAPIENS (HUMAN).//P39189

F-HEMBA1004711//ETS-RELATED PROTEIN 71 (ETS TRANSLATION VARIANT 2).//0.0027:148:30//HOMO SAPIENS (HUMAN).//000321

F-HEMBA1004725//CUTICLE COLLAGEN 2.//0.0051:41:41//CAENORHABDITIS ELEGANS.//P17656

F-HEMBA1004730//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//1.4e-22:210:37//HOMO SAPIENS (HUMAN).//P08547

F-HEMBA1004733//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//4.7e-07:50:62//HOMO SAPIENS (HUMAN).//P39188

F-HEMBA1004734//UBIQUITIN-CONJUGATING ENZYME E2-18 KD (EC 6.3.2.19) (UBIQUITIN- PROTEIN LIGASE) (UBIQUITIN CARRIER PROTEIN) (P42).//9.9e-39:143:52//ARABIDOPSIS THALIANA (MOUSE-EAR CRESS).//P42743

F-HEMBA1004736//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//4.1e-60:210:61//HOMO SAPIENS (HUMAN).//P08547

F-HEMBA1004748

F-HEMBA1004751//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//4.8e-20:88:63//HOMO SAPIENS (HUMAN).//P39188

F-HEMBA1004752//SKIN SECRETORY PROTEIN XP2 PRECURSOR (APEG PROTEIN).//0.0043:126:34//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P17437

F-HEMBA1004753//!!!! ALU SUBFAMILY SP WARNING ENTRY
!!!!//7.8e-28:47:78//HOMO SAPIENS (HUMAN).//P39193

F-HEMBA1004756//HYPOTHETICAL 53.3 KD PROTEIN IN HX 50

2896

T8-CAN1 INTERGENIC REGION.//0.22:77:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P39981

F-HEMBA1004758

F-HEMBA1004763//HYPOTHETICAL PROLINE-RICH PROTEIN (FRAGMENT).//1.1e-06:58:43//OWENIA FUSIFORMIS.//P21260

F-HEMBA1004768//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//4.7e-65:298:53//HOMO SAPIENS (HUMAN).//P08547

F-HEMBA1004770

F-HEMBA1004771

F-HEMBA1004776//GRANULIN 1.//0.78:28:42//CYPRIINUS CARPIO (COMMON CARP).//P81013

F-HEMBA1004778

F-HEMBA1004795//CDC4-LIKE PROTEIN (FRAGMENT).//6.9e-20:74:63//HOMO SAPIENS (HUMAN).//P50851

F-HEMBA1004803//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//1.4e-22:58:86//HOMO SAPIENS (HUMAN).//P08547

F-HEMBA1004806//HYPOTHETICAL 24.3 KD PROTEIN IN PS BH-RPL11 INTERGENIC REGION (ORF182).//0.72:75:33//CYANOPHORA PARADOXA.//P48324

F-HEMBA1004807

F-HEMBA1004816

F-HEMBA1004820//HEMOLYMPH TRYPSIN INHIBITOR A (BPI-TYPE) (FRAGMENT).//1.0:50:38//MANDUCA SEXTA (TOBACCO HAWKMOOTH) (TOBACCO HORNWORM).//P26226

F-HEMBA1004847//SIGNAL RECOGNITION PARTICLE 68 KD PROTEIN (SRP68).//3.0e-76:171:91//CANIS FAMILIARIS (DOG).//Q00004

F-HEMBA1004850//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//3.0e-05:64:43//BOS TAURUS (BOVINE).//P25508

F-HEMBA1004863//TOXIN C13S1C1 PRECURSOR.//0.38:52:30//DENDROASPIS ANGUSTICEPS (EASTERN GREEN MAMBA).//P18329

F-HEMBA1004864//TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN) (FRAGMENT).//0.89:24:50//HUMAN IMMUNODEFICIENCY VIRUS TYPE 1 (BH5 ISOLATE) (HIV-1).//P04612

F-HEMBA1004865

40 F-HEMBA1004880

F-HEMBA1004889//SMALL PROLINE-RICH PROTEIN II (SPR-II) (CLONE 174N).//0.66:23:47//HOMO SAPIENS (HUMAN).//P22532

F-HEMBA1004900

F-HEMBA1004909

F-HEMBA1004918//CHLOROPLAST 30S RIBOSOMAL PROTEIN S8 (FRAGMENT).//0.56:37:32//SPINACIA OLERACEA (SPINACH).//P09597

F-HEMBA1004923//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//3.5e-24:44:68//HOMO SAPIENS (HUMAN).//P39188

- 8
F-HEMBA1004929//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (A6L).//0.97:39:38//STRONGYLOCENTROTUS PURPURATUS (PURPLE SEA URCHIN).//P15997
F-HEMBA1004930//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//6.6e-15:64:59//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1004933//VASODILATOR-STIMULATED PHOSPHOPROTEIN (VASP).//0.34:58:41//HOMO SAPIENS (HUMAN).//P50552
F-HEMBA1004934
F-HEMBA1004944
F-HEMBA1004954//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 3 (EC 1.6.5.3).//0.58:78:30//PARAMECIUM TETRAURELIA.//P15579
F-HEMBA1004956//HYPOTHETICAL 18.8 KD PROTEIN (ORF 4).//0.98:57:31//PARAMECIUM TETRAURELIA.//P15605
F-HEMBA1004960//HYPOTHETICAL 12.6 KD PROTEIN (ORF J) (RETRON EC67).//1.0:58:27//ESCHERICHIA COLI.//P21324
F-HEMBA1004972
F-HEMBA1004973//SMALL PROLINE RICH PROTEIN II (SPR-II) (CLONE 930).//0.90:55:30//HOMO SAPIENS (HUMAN).//P22531
F-HEMBA1004977
F-HEMBA1004978
F-HEMBA1004980//MOTILIN PRECURSOR.//0.088:79:31//MACACA MULATTA (RHESUSMACAQUE).//O18811
F-HEMBA1004983//10 KD CHAPERONIN (PROTEIN CPN10) (PROTEIN GROES).//0.87:51:31//BUCHNERA APHIDICOLA.//Q59176
F-HEMBA1004995//MYOCYTE-SPECIFIC ENHANCER FACTOR 2B (SERUM RESPONSE FACTOR-LIKE PROTEIN 2) (XMEF2) (RSRFR2).//0.17:52:40//HOMO SAPIENS (HUMAN).//Q02080
F-HEMBA1005008//METALLOTHIONEIN (MT).//1.0:52:32//CRASSOSTREA VIRGINICA(EASTERN OYSTER).//P23038
F-HEMBA1005009//ACTIN.//3.5e-27:171:38//CANDIDA ALBICANS (YEAST).//P14235
F-HEMBA1005019//HYPOTHETICAL PROTEIN HI1222.//0.13:58:31//HAEMOPHILUS INFLUENZAE.//P44129
F-HEMBA1005029//P2Y PURINOCEPTOR 5 (P2Y5) (PURINERGIC RECEPTOR 5) (6H1).//0.76:72:31//GALLUS GALLUS (CHICKEN).//P32250
F-HEMBA1005035//HOMEBOX PROTEIN HB9.//0.0086:60:40//HOMO SAPIENS (HUMAN).//P50219
F-HEMBA1005039//SMALL PROLINE-RICH PROTEIN II (SPR-II) (CLONE 174N).//0.47:49:32//HOMO SAPIENS (HUMAN).//P22532
F-HEMBA1005047//RAS-RELATED PROTEIN RAB-24 (RAB-16).//1.5e-19:39:100//MUS MUSCULUS (MOUSE).//P35290
- 10
F-HEMBA1005050//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//0.074:34:44//BOS TAURUS (BOVINE).//P25508
F-HEMBA1005062
F-HEMBA1005066//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//2.1e-44:126:65//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1005075//SUPPRESSOR PROTEIN SRP40.//0.35:96:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P32583
F-HEMBA1005079//!!!! ALU SUBFAMILY SB2 WARNING ENTRY !!!!!/3.6e-20:75:64//HOMO SAPIENS (HUMAN).//P39191
F-HEMBA1005083//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//0.00015:72:34//BOS TAURUS (BOVINE).//P25508
F-HEMBA1005101//HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN 27C (HNRNP 48) (HRP48.1).//4.8e-10:176:25//DROSOPHILA MELANOGASTER (FRUIT FLY).//P48809
F-HEMBA1005113
F-HEMBA1005123//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/3.6e-24:99:60//HOMO SAPIENS (HUMAN).//P39194
F-HEMBA1005133//HYPOTHETICAL 13.5 KD PROTEIN IN MOB1-SGA1 INTERGENIC REGION.//0.11:22:54//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40490
F-HEMBA1005149//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!/3.7e-16:59:71//HOMO SAPIENS (HUMAN).//P39188
F-HEMBA1005152//GENOME POLYPROTEIN 2 [CONTAINS: HELPER COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); 70 KD PROTEIN].//1.0:77:27//BARLEY YELLOW MOSAIC VIRUS (JAPANESE STRAIN II-1) (BAYMV).//Q01207
F-HEMBA1005159//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 4L (EC 1.6.5.3).//0.40:53:33//APIS MELLIFERA (HONEYBEE).//P34859
F-HEMBA1005185//MYOSIN IB HEAVY CHAIN.//0.011:58:48//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P34092
F-HEMBA1005201//HYPOTHETICAL 56.6 KD PROTEIN C16C9.03 IN CHROMOSOME I.//3.9e-67:241:53//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09817
F-HEMBA1005202//SIGNAL RECOGNITION PARTICLE 68 KD PROTEIN (SRP68).//3.8e-124:257:95//CANIS FAMILIARIS (DOG).//Q00004
F-HEMBA1005206//CUTICLE COLLAGEN 1.//0.010:118:33//CAENORHABDITIS ELEGANS.//P08124
F-HEMBA1005219//PTB-ASSOCIATED SPLICING FACTOR (PSF).//0.99:85:40//HOMOSAPIENS (HUMAN).//P23246
F-HEMBA1005223//HYPOTHETICAL GENE 1.05 PROTEIN.//0.31:75:28//BACTERIOPHAGE T3.//P07715
F-HEMBA1005232//HYPOTHETICAL 7.8 KD PROTEIN.//0.99:48:29//VACCINIA VIRUS(STRAIN WR), AND VACCINIA V
- 20
Y !!!!!/3.6e-24:99:60//HOMO SAPIENS (HUMAN).//P39194
F-HEMBA1005133//HYPOTHETICAL 13.5 KD PROTEIN IN MOB1-SGA1 INTERGENIC REGION.//0.11:22:54//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40490
F-HEMBA1005149//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!/3.7e-16:59:71//HOMO SAPIENS (HUMAN).//P39188
F-HEMBA1005152//GENOME POLYPROTEIN 2 [CONTAINS: HELPER COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); 70 KD PROTEIN].//1.0:77:27//BARLEY YELLOW MOSAIC VIRUS (JAPANESE STRAIN II-1) (BAYMV).//Q01207
F-HEMBA1005159//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 4L (EC 1.6.5.3).//0.40:53:33//APIS MELLIFERA (HONEYBEE).//P34859
F-HEMBA1005185//MYOSIN IB HEAVY CHAIN.//0.011:58:48//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P34092
F-HEMBA1005201//HYPOTHETICAL 56.6 KD PROTEIN C16C9.03 IN CHROMOSOME I.//3.9e-67:241:53//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09817
F-HEMBA1005202//SIGNAL RECOGNITION PARTICLE 68 KD PROTEIN (SRP68).//3.8e-124:257:95//CANIS FAMILIARIS (DOG).//Q00004
F-HEMBA1005206//CUTICLE COLLAGEN 1.//0.010:118:33//CAENORHABDITIS ELEGANS.//P08124
F-HEMBA1005219//PTB-ASSOCIATED SPLICING FACTOR (PSF).//0.99:85:40//HOMOSAPIENS (HUMAN).//P23246
F-HEMBA1005223//HYPOTHETICAL GENE 1.05 PROTEIN.//0.31:75:28//BACTERIOPHAGE T3.//P07715
F-HEMBA1005232//HYPOTHETICAL 7.8 KD PROTEIN.//0.99:48:29//VACCINIA VIRUS(STRAIN WR), AND VACCINIA V
- 30
F-HEMBA1005152//GENOME POLYPROTEIN 2 [CONTAINS: HELPER COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); 70 KD PROTEIN].//1.0:77:27//BARLEY YELLOW MOSAIC VIRUS (JAPANESE STRAIN II-1) (BAYMV).//Q01207
F-HEMBA1005159//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 4L (EC 1.6.5.3).//0.40:53:33//APIS MELLIFERA (HONEYBEE).//P34859
F-HEMBA1005185//MYOSIN IB HEAVY CHAIN.//0.011:58:48//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P34092
F-HEMBA1005201//HYPOTHETICAL 56.6 KD PROTEIN C16C9.03 IN CHROMOSOME I.//3.9e-67:241:53//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09817
F-HEMBA1005202//SIGNAL RECOGNITION PARTICLE 68 KD PROTEIN (SRP68).//3.8e-124:257:95//CANIS FAMILIARIS (DOG).//Q00004
F-HEMBA1005206//CUTICLE COLLAGEN 1.//0.010:118:33//CAENORHABDITIS ELEGANS.//P08124
F-HEMBA1005219//PTB-ASSOCIATED SPLICING FACTOR (PSF).//0.99:85:40//HOMOSAPIENS (HUMAN).//P23246
F-HEMBA1005223//HYPOTHETICAL GENE 1.05 PROTEIN.//0.31:75:28//BACTERIOPHAGE T3.//P07715
F-HEMBA1005232//HYPOTHETICAL 7.8 KD PROTEIN.//0.99:48:29//VACCINIA VIRUS(STRAIN WR), AND VACCINIA V
- 40
F-HEMBA1005202//SIGNAL RECOGNITION PARTICLE 68 KD PROTEIN (SRP68).//3.8e-124:257:95//CANIS FAMILIARIS (DOG).//Q00004
F-HEMBA1005206//CUTICLE COLLAGEN 1.//0.010:118:33//CAENORHABDITIS ELEGANS.//P08124
F-HEMBA1005219//PTB-ASSOCIATED SPLICING FACTOR (PSF).//0.99:85:40//HOMOSAPIENS (HUMAN).//P23246
F-HEMBA1005223//HYPOTHETICAL GENE 1.05 PROTEIN.//0.31:75:28//BACTERIOPHAGE T3.//P07715
F-HEMBA1005232//HYPOTHETICAL 7.8 KD PROTEIN.//0.99:48:29//VACCINIA VIRUS(STRAIN WR), AND VACCINIA V
- 50
F-HEMBA1005232//HYPOTHETICAL 7.8 KD PROTEIN.//0.99:48:29//VACCINIA VIRUS(STRAIN WR), AND VACCINIA V

2899

2900

- IRUS (STRAIN COPENHAGEN). //P20544
 F-HEMBA1005241//!!!! ALU SUBFAMILY SP WARNING ENTR
 Y !!!!!/1.4e-28:138:55//HOMO SAPIENS (HUMAN). //P39
 193
 F-HEMBA1005244//SMALL PROLINE RICH PROTEIN II (SPR
 -II) (CLONE 930). //0.014:39:41//HOMO SAPIENS (HUMA
 N). //P22531
 F-HEMBA1005251//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
 4) (A6L). //0.55:15:46//DICENTRARCHUS LABRAX (EUROP
 EAN SEA BASS). //Q36362
 F-HEMBA1005252//EC PROTEIN HOMOLOG (ZINC-METALLOTH
 IONEIN CLASS II). //0.088:33:42//ZEA MAYS (MAIZE). /
 /P43401
 F-HEMBA1005274
 F-HEMBA1005275//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/0.96:42:45//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1005293//PROBABLE COATOMER BETA' SUBUNIT (B
 ETA'-COAT PROTEIN) (BETA'-COP). //0.55:98:30//CAENO
 RHABDITIS ELEGANS. //Q20168
 F-HEMBA1005296//MUCIN 2 PRECURSOR (INTESTINAL MUCI
 N 2). //0.095:75:34//HOMO SAPIENS (HUMAN). //Q02817
 F-HEMBA1005304//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/5.4e-33:103:74//HOMO SAPIENS (HUMAN). //P39
 189
 F-HEMBA1005311//PERIOD CLOCK PROTEIN (FRAGMENT). //
 0.99:45:31//DROSOPHILASALTANS (FRUIT FLY). //Q04536
 F-HEMBA1005314//HYPOTHETICAL 6.3 KD PROTEIN T19C3.
 3 IN CHROMOSOME III. //0.98:30:30//CAENORHABDITIS E
 LEGANS. //Q10009
 F-HEMBA1005315//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //1.1e-05:35:51//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1005318//OLFACTORY RECEPTOR-LIKE PROTEIN CO
 R8 (FRAGMENT). //0.57:44:38//GALLUS GALLUS (CHICKE
 N). //Q98913
 F-HEMBA1005331//IMMEDIATE-EARLY PROTEIN IE180. //0.
 57:106:33//PSEUDORABIES VIRUS (STRAIN INDIANA-FUNK
 HAUSER / BECKER) (PRV). //P11675
 F-HEMBA1005338//CARTILAGE MATRIX PROTEIN PRECURSOR
 (MATRILIN-1). //1.8e-55:199:59//GALLUS GALLUS (CHI
 CKEN). //P05099
 F-HEMBA1005353//CHLOROPLAST 30S RIBOSOMAL PROTEIN
 S17. //0.88:33:36//PORPHYRA PURPUREA. //P51305
 F-HEMBA1005359//ZINC FINGER PROTEIN 83 (ZINC FINGE
 R PROTEIN HPF1). //1.1e-68:255:48//HOMO SAPIENS (HU
 MAN). //P51522
 F-HEMBA1005367//ALPHA-AMYLASE INHIBITOR AAI. //1.0:
 25:40//AMARANTHUS HYPOCHONDRIACUS (PRINCE'S FEATHE
 R). //P80403
 F-HEMBA1005372
 F-HEMBA1005374//!!!! ALU SUBFAMILY SQ WARNING ENTR 50
 Y !!!!!/2.0e-34:92:75//HOMO SAPIENS (HUMAN). //P391
 94
 F-HEMBA1005382//APOLIPOPROTEIN C-II (APO-CII). //0.
 99:39:33//BOS TAURUS (BOVINE). //P19034
 F-HEMBA1005389//HYPOTHETICAL 70.0 KD PROTEIN IN DN
 AK 3' REGION (ORF4). //0.82:164:31//LACTOCOCCUS LACT
 IS (SUBSP. LACTIS) (STREPTOCOCCUS LACTIS). //P42377
 F-HEMBA1005394//HYPOTHETICAL 8.9 KD PROTEIN IN IE0
 -IE1 INTERGENIC REGION. //0.98:44:38//AUTOGRAPHA CA
 LIFORNICA NUCLEAR POLYHEDROSIS VIRUS (ACMPV). //P4
 1703
 F-HEMBA1005403//SPERM HISTONE P2 PRECURSOR (PROTAM
 INE MP2). //0.066:64:29//MUS MUSCULUS (MOUSE). //P07
 978
 F-HEMBA1005408//50S RIBOSOMAL PROTEIN L33. //0.77:3
 2:25//BACILLUS SUBTILIS. //Q06798
 F-HEMBA1005410//RETROVIRUS-RELATED POL POLYPROTEIN
 [CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); E
 NDONUCLEASE]. //0.0065:38:52//MUS MUSCULUS (MOUSE).
 20 //P11369
 F-HEMBA1005411//TOXIN S4C8. //0.16:46:28//DENDROASP
 IS JAMESONI KAIMOSAE (EASTERN JAMESON'S MAMBA). //P
 25683
 F-HEMBA1005423//CYCLIN-DEPENDENT KINASE 6 INHIBITO
 R (P18-INK6) (CYCLIN-DEPENDENT KINASE 4 INHIBITOR
 C) (P18-INK4C). //4.3e-09:29:96//HOMO SAPIENS (HUMA
 N). //P42773
 F-HEMBA1005426//TOXIN C10S2C2. //0.99:49:34//DENDRO
 ASPIS ANGUSTICEPS (EASTERN GREEN MAMBA). //P25684
 F-HEMBA1005443//!!!! ALU SUBFAMILY J WARNING ENTRY 30
 !!!!!/2.9e-16:78:60//HOMO SAPIENS (HUMAN). //P3918
 8
 F-HEMBA1005447//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
 4) (A6L). //0.99:57:31//DASYPUS NOVENMCINCTUS (NINE-
 BANDED ARMADILLO). //021329
 F-HEMBA1005468//NADH-UBIQUINONE OXIDOREDUCTASE CHA
 IN 1 (EC 1.6.5.3) (FRAGMENTS). //0.68:41:31//ARTEMI
 A SALINA (BRINE SHRIMP). //P19040
 F-HEMBA1005469
 F-HEMBA1005472//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //1.5e-39:142:70//HOMO SAPIENS (HUMAN). //P08547
 F-HEMBA1005474//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!/5.8e-10:44:68//HOMO SAPIENS (HUMAN). //P391
 94
 F-HEMBA1005475//U1 SMALL NUCLEAR RIBONUCLEOPROTEIN
 70 KD (U1 SNRNP 70 KD) (SNRNP70). //9.2e-14:179:33/
 /HOMO SAPIENS (HUMAN). //P08621
 F-HEMBA1005497
 F-HEMBA1005500//60S RIBOSOMAL PROTEIN L37. //0.11:5
 3:33//SCHISTOSOMA MANSONI (BLOOD FLUKE). //044125

F-HEMBA1005506

F-HEMBA1005508

F-HEMBA1005511//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/3.5e-30:92:73//HOMO SAPIENS (HUMAN).//P391
94

F-HEMBA1005513//MALES-ABSENT ON THE FIRST PROTEIN
(EC 2.3.1.-).//2.0e-39:95:61//DROSOPHILA MELANOGAS
TER (FRUIT FLY).//002193

F-HEMBA1005517//PROLINE-RICH PROTEIN MP-2 PRECURSO
R.//2.1e-06:56:44//MUSMUSCULUS (MOUSE).//P05142

F-HEMBA1005518//COLLAGEN ALPHA 1(I) CHAIN (FRAGMEN
TS).//5.8e-05:192:33//BOS TAURUS (BOVINE).//P02453

F-HEMBA1005520//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/2.0e-18:87:57//HOMO SAPIENS (HUMAN).//P3918
8

F-HEMBA1005526//!!!! ALU SUBFAMILY SB2 WARNING ENT
RY !!!!!/5.1e-22:77:54//HOMO SAPIENS (HUMAN).//P39
191

F-HEMBA1005528//CCR4-ASSOCIATED FACTOR 1 (CAF1).//
1.2e-81:157:98//MUS MUSCULUS (MOUSE).//Q60809

F-HEMBA1005530//POLLEN ALLERGEN AMB P 5-A PRECURSO
R (AMB P V-A).//0.98:19:47//AMBROSIA PSILOSTACHYA
(WESTERN RAGWEED).//P43174

F-HEMBA1005548//TRANSCRIPTION FACTOR MAF1.//1.4e-7
2:137:97//RATTUS NORVEGICUS (RAT).//P54842

F-HEMBA1005552//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/1.7e-29:47:78//HOMO SAPIENS (HUMAN).//P391
93

F-HEMBA1005558//HYPOTHETICAL 25.6 KD PROTEIN IN AB
F2-CHL12 INTERGENIC REGION.//1.6e-20:202:30//SACCH
AROMYCES CEREVISIAE (BAKER'S YEAST).//Q04272F-HEMB
A1005568

F-HEMBA1005570//NADH-UBIQUINONE OXIDOREDUCTASE CHA
IN 6 (EC 1.6.5.3).//1.0:80:31//CAENORHABDITIS ELEG
ANS.//P24885

F-HEMBA1005576//TRANSMEMBRANE PROTEIN SEX PRECURSO
R.//8.5e-58:152:75//HOMO SAPIENS (HUMAN).//P51805

F-HEMBA1005577//KERATIN, HIGH-SULFUR MATRIX PROTEI
N, B2A.//0.98:57:36//OVIS ARIES (SHEEP).//P02438

F-HEMBA1005581//SLIT PROTEIN PRECURSOR.//1.1e-62:2
54:41//DROSOPHILA MELANOGASTER (FRUIT FLY).//P2401
4

F-HEMBA1005582//DYNACTIN, 150 KD ISOFORM (150 KD D
YNEIN-ASSOCIATED POLYPEPTIDE) (DP-150) (DAP-150)
(P150-GLUED).//0.0091:189:29//RATTUS NORVEGICUS (R
AT).//P28023

F-HEMBA1005583//HYPOTHETICAL 41.2 KD PROTEIN IN CP
S REGION (ORF7).//0.83:119:23//KLEBSIELLA PNEUMONI
AE.//Q48453

F-HEMBA1005588//!!!! ALU SUBFAMILY J WARNING ENTRY 50

!!!!/1.9e-17:108:53//HOMO SAPIENS (HUMAN).//P391
88

F-HEMBA1005593//SMALL PROLINE-RICH PROTEIN II (SPR
-II) (CLONE 174N).//0.23:24:54//HOMO SAPIENS (HUMA
N).//P22532

F-HEMBA1005595//DYNEIN HEAVY CHAIN, CYTOSOLIC (DYH
C).//2.7e-39:257:39//DICTYOSTELIUM DISCOIDEUM (SLI
ME MOLD).//P34036

F-HEMBA1005606

F-HEMBA1005609//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!/3.2e-20:27:96//HOMO SAPIENS (HUMAN).//P391
92

F-HEMBA1005616//LATE CONTROL GENE B PROTEIN (GPB).
//0.48:51:33//BACTERIOPHAGE 186.//P08711

F-HEMBA1005621//MITOTIC MAD2 PROTEIN.//1.2e-06:13
7:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P
40958

F-HEMBA1005627//HYPOTHETICAL 17.1 KD PROTEIN IN PU
R5 3' REGION.//0.18:100:32//SACCHAROMYCES CEREVISIA
E (BAKER'S YEAST).//P38898

F-HEMBA1005631

F-HEMBA1005632//HYPOTHETICAL 7.4 KD PROTEIN.//0.3
2:59:32//VACCINIA VIRUS(STRAIN WR).//P04309

F-HEMBA1005634//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/2.6e-14:93:58//HOMO SAPIENS (HUMAN).//P3918
8

F-HEMBA1005666//HYPOTHETICAL PROTEIN KIAA0129.//2.
1e-05:126:25//HOMO SAPIENS (HUMAN).//Q14142

F-HEMBA1005670

30 F-HEMBA1005679//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//2.2e-08:40:72//HOMO SAPIENS (HUMAN).//P08547

F-HEMBA1005680//SMALL PROLINE-RICH PROTEIN 2-1.//
0.015:19:47//HOMO SAPIENS (HUMAN).//P35326

F-HEMBA1005685

F-HEMBA1005699//EPHRIIN-B3 PRECURSOR (EPH-RELATED R
ECEPTOR TYROSINE KINASE LIGAND 8) (LERK-8) (EPH-RE
LATED RECEPTOR TRANSMEMBRANE LIGAND ELK-L3).//4.2e
-38:98:81//HOMO SAPIENS (HUMAN).//Q15768

F-HEMBA1005705//PROTEIN Q300.//0.11:23:56//MUS MUS
CULUS (MOUSE).//Q02722F-HEMBA1005717

F-HEMBA1005732//BACTENECIN 7 PRECURSOR (BAC7).//0.
22:55:41//OVIS ARIES (SHEEP).//P50415

F-HEMBA1005737//CALCINEURIN B SUBUNIT (PROTEIN PHO
SPHATASE 2B REGULATORYSUBUNIT).//4.5e-18:167:34//S
ACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P25296

F-HEMBA1005746

F-HEMBA1005755//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//7.4e-30:69:65//HOMO SAPIENS (HUMAN).//P08547

F-HEMBA1005765//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/2.8e-19:60:63//HOMO SAPIENS (HUMAN).//P391

94

F-HEMBA1005780//METALLOTHIONEIN-I (MT-1).//1.0:31:38//COLUMBA LIVIA (DOMESTIC PIGEON).//P15786

F-HEMBA1005813

F-HEMBA1005815//CALPAIN, LARGE [CATALYTIC] SUBUNIT (EC 3.4.22.17) (CALCIUM- ACTIVATED NEUTRAL PROTEINASE) (CANP) (MU/M-TYPE).//1.0e-23:200:31//GALLUS GALLUS (CHICKEN).//P00789

F-HEMBA1005822//PROTEIN Q300.//0.0016:21:80//MUS MUSCULUS (MOUSE).//Q02722

F-HEMBA1005829//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/9.6e-33:96:73//HOMO SAPIENS (HUMAN).//P391

94

F-HEMBA1005834//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//1.6e-22:103:46//NYCTICEBUS COUCANG (SLOW LORIS).//P08548

F-HEMBA1005852//PROLINE-RICH PROTEIN MP-3 (FRAGMENT).//8.8e-06:95:35//MUS MUSCULUS (MOUSE).//P05143

F-HEMBA1005853//HYPOTHETICAL PROTEIN MJ0647.//0.39:28:39//METHANOCOCCUS JANNASCHII.//Q58063

F-HEMBA1005884

F-HEMBA1005891//HYPOTHETICAL PROTEIN MTH137.//0.95:51:27//METHANOBACTERIUM THERMOAUTOTROPHICUM.//026240

F-HEMBA1005894//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!/2.6e-29:81:71//HOMO SAPIENS (HUMAN).//P391

95

F-HEMBA1005909//HYPOTHETICAL 8.2 KD PROTEIN B0353.1 IN CHROMOSOME III.//0.98:19:52//CAENORHABDITIS ELEGANS.//Q10958

F-HEMBA1005911//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!/1.9e-27:86:70//HOMO SAPIENS (HUMAN).//P3918

8

F-HEMBA1005921//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/1.3e-38:99:81//HOMO SAPIENS (HUMAN).//P391

94

F-HEMBA1005931//ZINC FINGER PROTEIN 83 (ZINC FINGER PROTEIN HPF1).//2.3e-17:76:51//HOMO SAPIENS (HUMAN).//P51522

F-HEMBA1005934//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!/0.024:54:40//HOMO SAPIENS (HUMAN).//P39189

F-HEMBA1005962

F-HEMBA1005963//COATOMER GAMMA SUBUNIT (GAMMA-COAT PROTEIN) (GAMMA-COP).//1.7e-32:89:79//BOS TAURUS (BOVINE).//P53620

F-HEMBA1005990//HYPOTHETICAL BHLF1 PROTEIN.//3.0e-09:180:36//EPSTEIN-BARR VIRUS (STRAIN B95-8) (HUMAN HERPESVIRUS 4).//P03181

F-HEMBA1005991//HYPOTHETICAL PROTEIN KIAA0032.//3.0e-17:107:43//HOMO SAPIENS (HUMAN).//Q15034

F-HEMBA1005999

F-HEMBA1006002

F-HEMBA1006005//CORNIFIN B (SMALL PROLINE-RICH PROTEIN 1B) (SPR1B) (SPR1B).//0.0017:45:44//MUS MUSCULUS (MOUSE).//Q62267

F-HEMBA1006031//BASIC PROLINE-RICH PEPTIDE 1B-1.//0.00016:84:39//HOMO SAPIENS (HUMAN).//P04281

F-HEMBA1006035//DNAK PROTEIN 1 (HEAT SHOCK PROTEIN 70) (HSP70).//0.43:100:27//SYNECHOCYSTIS SP. (STRAIN PCC 6803).//Q55154

10

F-HEMBA1006036//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/6.2e-64:150:74//HOMO SAPIENS (HUMAN).//P39

194

F-HEMBA1006042

F-HEMBA1006067//METALLOTHIONEIN A (MT-A).//0.86:34:41//THERMARCUS CERBERUS.//P52721

F-HEMBA1006081

F-HEMBA1006090//SODIUM/GLUCOSE COTRANSPORTER 3 (NA (+)/GLUCOSE COTRANSPORTER 3) (LOW AFFINITY SODIUM-GLUCOSE COTRANSPORTER).//0.87:35:54//SUS SCROFA (PIG).//P31636

20

F-HEMBA1006091//EARLY NODULIN 20 PRECURSOR (N-20).//0.027:87:32//MEDICAGO TRUNCATULA (BARREL MEDIC).//P93329

F-HEMBA1006100//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!/8.1e-09:58:60//HOMO SAPIENS (HUMAN).//P391

95

F-HEMBA1006108//HYPOTHETICAL 56.6 KD PROTEIN IN UR E2-SSU72 INTERGENIC REGION.//5.6e-16:88:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53867

30

F-HEMBA1006121//HOMEBOX PROTEIN CDX-1 (CAUDAL-TYPE HOMEBOX PROTEIN 1).//3.4e-05:106:37//HOMO SAPIENS (HUMAN).//P47902

F-HEMBA1006124//50S RIBOSOMAL PROTEIN L33.//1.0:12:83//BACILLUS STEAROTHERMOPHILUS.//P23375

F-HEMBA1006130//SEL-10 PROTEIN.//7.7e-05:129:28//CAENORHABDITIS ELEGANS.//Q93794

F-HEMBA1006138//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/7.8e-13:41:73//HOMO SAPIENS (HUMAN).//P391

94

F-HEMBA1006142//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!/2.3e-39:101:77//HOMO SAPIENS (HUMAN).//P39

192

F-HEMBA1006155//GENE 33 POLYPEPTIDE.//0.21:70:31//RATTUS NORVEGICUS (RAT).//P05432

F-HEMBA1006158

F-HEMBA1006173//PROTEIN-TYROSINE PHOSPHATASE STRIATUM-ENRICHED (EC 3.1.3.48) (STEP) (NEURAL-SPECIFIC PROTEIN-TYROSINE PHOSPHATASE) (FRAGMENT).//0.017:

50

20:95//HOMO SAPIENS (HUMAN).//P54829

2905

2906

F-HEMBA1006182//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/0.37:31:61//HOMO SAPIENS (HUMAN).//P39188
 F-HEMBA1006198//HOMEODOMAIN PROTEIN HOX-B3 (HOX-2.7)
 (MH-23).//0.85:61:29//MUS MUSCULUS (MOUSE).//P0902
 6
 F-HEMBA1006235//50S RIBOSOMAL PROTEIN L33.//1.0:2
 6:38//AQUIFEX AEOLICUS.//067756
 F-HEMBA1006248//MALE SPECIFIC SPERM PROTEIN MST84D
 B.//0.0041:64:37//DROSOPHILA MELANOGASTER (FRUIT F
 LY).//Q01643
 F-HEMBA1006252//BOWMAN-BIRK TYPE PROTEINASE INHIBI
 TOR DE-3.//1.0:22:40//DOLICHOS AXILLARIS (MACROTYL
 OMA AXILLARE).//P01057
 F-HEMBA1006253//DISINTEGRIN ERISTICOPHIN (PLATELET
 AGGREGATION ACTIVATION INHIBITOR).//0.95:19:47//E
 RISTICOPHIS MACMAHONI (LEAF-NOSED VIPER).//P22826
 F-HEMBA1006259
 F-HEMBA1006268//!!!! ALU SUBFAMILY SC WARNING ENTR
 Y !!!!!/7.0e-05:32:65//HOMO SAPIENS (HUMAN).//P391
 92
 F-HEMBA1006272//RETROVIRUS-RELATED GAG POLYPROTEIN
 (VERSION 2).//4.8e-112:248:78//HOMO SAPIENS (HUMA
 N).//P10264
 F-HEMBA1006278//POLY(A) POLYMERASE (EC 2.7.7.19)
 (PAP) (POLYNUCLEOTIDE ADENYLYLTRANSFERASE) (FRAGME
 NT).//2.5e-71:164:75//HOMO SAPIENS (HUMAN).//P5100
 3
 F-HEMBA1006283//50S RIBOSOMAL PROTEIN L32.//0.81:2
 7:44//THERMUS AQUATICUS (SUBSP. THERMOPHILUS).//P8
 0339
 F-HEMBA1006284//CUTICLE COLLAGEN 2.//0.36:42:40//C
 AENORHABDITIS ELEGANS.//P17656
 F-HEMBA1006291//HYPOTHETICAL 43.3 KD PROTEIN IN EV
 GS-CLK INTERGENIC REGION.//2.4e-37:143:31//ESCHERI
 CHIA COLI.//P76518
 F-HEMBA1006293//MYELIN-OLIGODENDROCYTE GLYCOPROTEI
 N PRECURSOR.//0.20:134:29//RATTUS NORVEGICUS (RA
 T).//Q63345
 F-HEMBA1006309//HYPOTHETICAL 54.2 KD PROTEIN IN ER
 P5-ORC6 INTERGENIC REGION.//2.1e-43:187:48//SACCHA
 ROMYCES CEREVISIAE (BAKER'S YEAST).//P38821
 F-HEMBA1006310//SIGNAL TRANSDUCER CD24 PRECURSOR
 (HEAT STABLE ANTIGEN) (HSA) (NECTADRIN).//0.71:46:
 39//RATTUS NORVEGICUS (RAT).//Q07490
 F-HEMBA1006328//RNA POLYMERASE ALPHA SUBUNIT (EC
 2.7.7.48) (NUCLEOCAPSIDPHOSPHOPROTEIN).//0.44:141:
 24//HUMAN PARAINFLUENZA 1 VIRUS (STRAIN CI-5/73)./
 /P32531
 F-HEMBA1006334//HYPOTHETICAL TRANSCRIPTIONAL REGUL
 ATOR AF1627.//0.98:26:46//ARCHAEOGLOBUS FULGIDUS./

/028646
 F-HEMBA1006344//EZRIN (P81) (CYTOVILLIN) (VILLIN-
 2).//8.8e-08:91:36//MUSMUSCULUS (MOUSE).//P26040
 F-HEMBA1006347//MALES-ABSENT ON THE FIRST PROTEIN
 (EC 2.3.1.-).//9.1e-48:149:50//DROSOPHILA MELANOCA
 STER (FRUIT FLY).//002193
 F-HEMBA1006349//METALLOTHIONEIN-LIKE PROTEIN 1.//
 0.015:59:33//CASUARINAGLAUCA (SWAMP OAK).//Q39511
 F-HEMBA1006359//ZINC FINGER PROTEIN 43 (ZINC PROTE
 10 IN HTF6).//6.8e-96:261:66//HOMO SAPIENS (HUMAN).//
 P28160
 F-HEMBA1006364//PUTATIVE ENDONUCLEASE C1F12.06C (E
 C 3.1.-).//0.97:60:35//SCHIZOSACCHAROMYCES POMBE
 (FISSION YEAST).//Q10348
 F-HEMBA1006377//EARLY NODULIN 20 PRECURSOR (N-20).
 //0.00023:110:35//MEDICAGO TRUNCATULA (BARREL MEDI
 C).//P93329
 F-HEMBA1006380
 F-HEMBA1006381//METALLOTHIONEIN-II.//1.0:26:38//CA
 20 NDIDA GLABRATA (YEAST) (TORULOPSIS GLABRATA).//P151
 14
 F-HEMBA1006398//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG.//3.3e-26:123:52//HOMO SAPIENS (HUMAN).//P08547
 F-HEMBA1006416
 F-HEMBA1006419//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/1.2e-24:102:50//HOMO SAPIENS (HUMAN).//P39
 189
 F-HEMBA1006421//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/3.1e-21:101:57//HOMO SAPIENS (HUMAN).//P391
 30 88
 F-HEMBA1006424//HYPOTHETICAL PROTEIN IORF1.//0.85:
 55:30//BOVINE CORONAVIRUS (STRAIN MEBUS), AND BOVI
 NE CORONAVIRUS (STRAIN QUEBEC).//P22053
 F-HEMBA1006426//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!/1.8e-36:78:74//HOMO SAPIENS (HUMAN).//P391
 95
 F-HEMBA1006438//HYPOTHETICAL 8.1 KD PROTEIN (ORF6
 5).//1.0:38:36//GUILLARDIA THETA (CRYPTOMONAS PH
 I).//078421
 F-HEMBA1006445//RAS-LIKE PROTEIN 3.//1.9e-06:40:47
 //RHIZOMUCOR RACEMOSUS(MUCOR CIRCINELLOIDES F. LUS
 ITANICUS).//P22280
 F-HEMBA1006446
 F-HEMBA1006461//!!!! ALU SUBFAMILY SC WARNING ENTR
 Y !!!!!/4.1e-18:68:67//HOMO SAPIENS (HUMAN).//P391
 92
 F-HEMBA1006467
 F-HEMBA1006471
 F-HEMBA1006474//40 KD PROTEIN.//1.1e-37:231:38//BO
 50 RNA DISEASE VIRUS (BDV).//Q01552

2907

F-HEMBA1006483//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!/6.1e-38:77:74//HOMO SAPIENS (HUMAN).//P391
92
F-HEMBA1006485//HYPOTHETICAL 9.3 KD PROTEIN IN NAD
3-NAD7 INTERGENIC REGION (ORF 79).//0.91:30:40//MA
RCHANTIA POLYMORPHA (LIVERWORT).//P38465
F-HEMBA1006486//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//1.1e-12:78:51//HOMO SAPIENS (HUMAN).//P08547
F-HEMBA1006489//FUN34 PROTEIN.//0.94:58:37//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P32907
F-HEMBA1006492//NADH-UBIQUINONE OXIDOREDUCTASE MWF
E SUBUNIT (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-MWF
E) (CI-MWFE).//0.87:44:36//HOMO SAPIENS (HUMAN).//
015239
F-HEMBA1006494//FERREDOXIN-LIKE PROTEIN IN NIF REG
ION.//0.11:46:26//RHIZOBIUM LEGUMINOSARUM (BIOVAR
TRIFOLIUM).//P42711
F-HEMBA1006497
F-HEMBA1006502//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/0.15:26:73//HOMO SAPIENS (HUMAN).//P39188
F-HEMBA1006507//DIAPHANOUS PROTEIN.//0.0055:129:28
//DROSOPHILA MELANOCASTER (FRUIT FLY).//P48608
F-HEMBA1006521//3-OXOACYL- [ACYL-CARRIER PROTEIN] R
EDUCTASE (EC 1.1.1.100) (3-KETOACYL- ACYL CARRIER
PROTEIN REDUCTASE).//1.1e-32:177:41//ESCHERICHIA C
OLI.//P25716
F-HEMBA1006530//NADH-UBIQUINONE OXIDOREDUCTASE CHA
IN 5 (EC 1.6.5.3) (FRAGMENT).//0.052:84:26//LEISHM
ANIA TARENTOLAE (SAUROLEISHMANIA TARENTOLAE).//P15
583
F-HEMBA1006535//INHIBITOR OF APOPTOSIS PROTEIN 1
(MIAP1) (MIAP-1).//6.6e-05:53:39//MUS MUSCULUS (MO
USE).//008863
F-HEMBA1006540//PRESYNAPTIC PROTEIN SAP97 (SYNAPSE
-ASSOCIATED PROTEIN 97) (DISCS, LARGE HOMOLOG 1)./
/2.1e-07:206:23//RATTUS NORVEGICUS (RAT).//Q62696
F-HEMBA1006546//PROBABLE E5 PROTEIN.//0.11:70:32//
HUMAN PAPILLOMAVIRUS TYPE 51.//P26553
F-HEMBA1006559//SUPPRESSOR PROTEIN SRP40.//0.015:2
21:20//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).// 40
P32583
F-HEMBA1006562//SALIVARY PROLINE-RICH PROTEIN PO P
RECURSOR (ALLELE S).//1.5e-07:122:33//HOMO SAPIENS
(HUMAN).//P10163
F-HEMBA1006566//CELL DIVISION PROTEIN KINASE 2 (EC
2.7.1.-) (CDC2 HOMOLOG EG1 PROTEIN KINASE).//0.6
3:53:37//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P2
3437
F-HEMBA1006569//COLLAGEN ALPHA 2(I) CHAIN (FRAGMEN
T).//4.4e-06:88:39//BOS TAURUS (BOVINE).//P02465 50

2908

F-HEMBA1006579
F-HEMBA1006583//PROLINE-RICH PROTEIN MP-2 PRECURSO
R.//0.011:61:40//MUS MUSCULUS (MOUSE).//P05142
F-HEMBA1006595//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/5.6e-34:93:77//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBA1006597//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/1.9e-26:75:74//HOMO SAPIENS (HUMAN).//P391
95
F-HEMBA1006612//SUPPRESSOR PROTEIN SRP40.//0.026:2
21:22//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//
P32583
F-HEMBA1006617//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/6.6e-20:73:63//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1006624//HYPOTHETICAL 41.9 KD PROTEIN IN SD
S3-TSH1 INTERGENIC REGION.//2.6e-31:209:44//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P40506F-HEMBA
1006631//HYPOTHETICAL 62.8 KD PROTEIN IN TAF145-YO
R1 INTERGENIC REGION.//1.5e-15:131:41//SACCHAROMYC
ES CEREVISIAE (BAKER'S YEAST).//P53331
F-HEMBA1006635
F-HEMBA1006639//POLYADENYLATE-BINDING PROTEIN 1 (P
OLY(A) BINDING PROTEIN1) (PABP 1).//2.2e-11:48:75/
/MUS MUSCULUS (MOUSE).//P29341
F-HEMBA1006643//LONG NEUROTOXIN CR1 PRECURSOR (KAP
PA NEUROTOXIN).//0.28:48:27//BUNGARUS MULTICINCTUS
(MANY-BANDED KRAIT).//P15817
F-HEMBA1006648//ZINC FINGER PROTEIN 12 (ZINC FINGE
R PROTEIN KOX3) (FRAGMENT).//0.26:17:47//HOMO SAPI
ENS (HUMAN).//P17014
F-HEMBA1006652//60S RIBOSOMAL PROTEIN L7.//2.4e-4
4:206:47//MUS MUSCULUS(MOUSE).//P14148
F-HEMBA1006653
F-HEMBA1006659
F-HEMBA1006665//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//0.018:43:58//HOMOSAPIENS (HUMAN).//P08547
F-HEMBA1006674//TRANSCRIPTION INITIATION FACTOR TF
IID 135 KD SUBUNIT (TAFII-135) (TAFII135) (TAFII-1
30) (TAFII130).//2.9e-05:154:33//HOMO SAPIENS (HUM
AN).//000268
F-HEMBA1006676//HYPOTHETICAL PROLINE-RICH PROTEIN
(FRAGMENT).//3.6e-09:52:51//OWENIA FUSIFORMIS.//P2
1260
F-HEMBA1006682
F-HEMBA1006695//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/2.1e-06:35:65//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBA1006696
F-HEMBA1006708//HYPOTHETICAL 46.4 KD TRP-ASP REPEA

2909

TS CONTAINING PROTEININ PMC1-TFG2 INTERGENIC REGION. //3.4e-19:104:45//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53196

F-HEMBA1006709//RETINOIC ACID RECEPTOR RXR-BETA. //0.24:111:36//HOMO SAPIENS (HUMAN). //P28702

F-HEMBA1006717

F-HEMBA1006737//ANKYRIN, BRAIN VARIANT 2 (ANKYRIN B) (ANKYRIN, NONERYTHROID) (FRAGMENT). //5.8e-09:111:40//HOMO SAPIENS (HUMAN). //Q01485

F-HEMBA1006744//!!!! ALU SUBFAMILY SB2 WARNING ENTRY !!!!!//1.8e-32:84:78//HOMO SAPIENS (HUMAN). //P39191

F-HEMBA1006754//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //1.3e-75:220:62//HOMO SAPIENS (HUMAN). //P08547

F-HEMBA1006758//VASCULAR ENDOTHELIAL-CADHERIN PRECURSOR (VE-CADHERIN) (CADHERIN-5) (7B4 ANTIGEN) (CD144 ANTIGEN). //0.024:110:29//HOMO SAPIENS (HUMAN). //P33151

F-HEMBA1006767

F-HEMBA1006779//MITOCHONDRIAL RIBOSOMAL PROTEIN S12. //0.67:19:42//LEISHMANIA TARENTOLAE (SAUROLEISHMANIA TARENTOLAE). //Q34940

F-HEMBA1006780

F-HEMBA1006789//PROLINE-RICH PROTEIN MP-3 (FRAGMENT). //0.056:98:30//MUS MUSCULUS (MOUSE). //P05143

F-HEMBA1006795//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG. //2.9e-11:143:30//NYCTICEBUS COUCANG (SLOW LORIS). //P08548

F-HEMBA1006796//WISKOTT-ALDRICH SYNDROME PROTEIN HOMOLOG (WASP). //0.16:38:42//MUS MUSCULUS (MOUSE). //P70315

F-HEMBA1006807//HYPOTHETICAL 46.4 KD PROTEIN T16H12.5 IN CHROMOSOME III. //4.4e-75:184:77//CAENORHABDITIS ELEGANS. //P34568

F-HEMBA1006821//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//0.011:20:85//HOMO SAPIENS (HUMAN). //P39194

F-HEMBA1006824//PROTEIN B11. //0.44:27:44//VACCINIA VIRUS (STRAIN WR). //Q01229

F-HEMBA1006832//HYPOTHETICAL 34.6 KD PROTEIN C13G5.2 IN CHROMOSOME III. //1.0:46:36//CAENORHABDITIS ELEGANS. //P34327

F-HEMBA1006849

F-HEMBA1006865//ACROSIN INHIBITORS IIA AND IIB (BUSH-II). //1.0:41:31//BOS TAURUS (BOVINE). //P01001

F-HEMBA1006877//OXYSTEROL-BINDING PROTEIN. //3.7e-26:239:36//ORYZOLAGUSCUNICULUS (RABBIT). //P16258

F-HEMBA1006885//HYPOTHETICAL 27.2 KD PROTEIN F09E5.8 IN CHROMOSOME II. //4.5e-38:185:43//CAENORHABDITIS ELEGANS. //P52057

F-HEMBA1006900

2910

F-HEMBA1006914//UBIQUITIN-ACTIVATING ENZYME E1-LIKE (POLYMERASE-INTERACTING PROTEIN 2). //5.2e-27:269:33//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P52488

F-HEMBA1006921//CYTOTOXIN 3 (COMPONENT 3.20). //0.99:32:37//NAJA MELANOLEUCA (FOREST COBRA) (BLACK-LIPPED COBRA). //P01473

F-HEMBA1006926//ATROPHIN-1 (DENTATORUBRAL-PALLIDOLUYSIAN ATROPHY PROTEIN). //0.0024:148:33//RATTUS NORVEGICUS (RAT). //P54258

F-HEMBA1006929//HYPOTHETICAL PROTEIN MJ0525. //0.95:35:20//METHANOCOCCUSJANNASCHII. //Q57945

F-HEMBA1006936//SALIVARY ACIDIC PROLINE-RICH PHOSPHOPROTEIN 1/2 PRECURSOR (PRP-1 / PRP-3) (PRP-2 / PRP-4) (PIF-F / PIF-S) (PROTEIN A / PROTEIN C) [CONTAINS: PEPTIDE P-C]. //0.074:116:31//HOMO SAPIENS (HUMAN). //P02810

F-HEMBA1006938

F-HEMBA1006941//THIOREDOXIN H-TYPE 1 (TRX-H1). //2.1e-13:90:33//NICOTIANATABACUM (COMMON TOBACCO). //P29449

F-HEMBA1006949

F-HEMBA1006973//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS). //0.75:29:55//BOS TAURUS (BOVINE). //P25508

F-HEMBA1006976//CMP-N-ACETYLNEURAMINATE-BETA-GALACTOSAMIDE-ALPHA-2,3-SIALYLTRANSFERASE (EC 2.4.99.-) (BETA-GALACTOSIDE ALPHA-2,3-SIALYLTRANSFERASE) (ST3GALIII) (ALPHA 2,3-ST) (GAL-NAC6S) (STZ) (SIAT4-C) (SAT-3) (ST-4). //3.9e-108:117:95//HOMO SAPIENS (HUMAN). //Q11206

F-HEMBA1006993

F-HEMBA1006996//HYPOTHETICAL 8.7 KD PROTEIN IN RPL22-RPL23 INTERGENIC REGION (ORF70). //0.12:51:33//ASTASIA LONGA (EUGLENOPHYCEAN ALGA). //P34779

F-HEMBA1007002//PLATELET GLYCOPROTEIN IX PRECURSOR (GPIX) (CD42A). //0.00096:60:33//HOMO SAPIENS (HUMAN). //P14770

F-HEMBA1007017//HYPOTHETICAL 7.2 KD PROTEIN IN CYADAPF INTERGENIC REGION. //1.0:25:56//ESCHERICHIA COLI. //P39166

F-HEMBA1007018//DYNEIN LIGHT INTERMEDIATE CHAIN 1, CYTOSOLIC (LIC57/59) (DYNEIN LIGHT CHAIN A) (DLC-A). //8.5e-120:278:80//GALLUS GALLUS (CHICKEN). //Q90828

F-HEMBA1007045//SKIN SECRETORY PROTEIN XP2 PRECURSOR (APEG PROTEIN). //2.1e-12:158:29//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P17437

F-HEMBA1007051

F-HEMBA1007052//60S RIBOSOMAL PROTEIN L37-B (L35) (YP55). //0.94:37:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53196

50

- ER'S YEAST). //P51402
 F-HEMBA1007062//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3 4) (A6L). //0.93:55:29//RHINOCEROS UNICORNIS (GREATER INDIAN RHINOCEROS). //Q96063
 F-HEMBA1007066//ECLOSION HORMONE PRECURSOR (ECDYSIS ACTIVATOR) (EH). //0.58:49:38//BOMBYX MORI (SILK MOTH). //P25331
 F-HEMBA1007073//PUTATIVE SMALL MEMBRANE PROTEIN (ORF 4). //0.86:46:34//CANINE ENTERIC CORONAVIRUS (STRAIN INSAVC-1) (CCV). //P36696
 F-HEMBA1007078//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!/8.6e-29:56:67//HOMO SAPIENS (HUMAN). //P39193
 F-HEMBA1007080//SKIN SECRETORY PROTEIN XP2 PRECURSOR (APEG PROTEIN). //0.028:122:30//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P17437
 F-HEMBA1007085//RTOA PROTEIN (RATIO-A). //7.4e-11:221:31//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD). //P54681
 F-HEMBA1007087//HYPOTHETICAL PROTEIN MJ0162. //3.3e-29:173:36//METHANOCOCCUS JANNASCHII. //Q57626
 F-HEMBA1007112
 F-HEMBA1007113
 F-HEMBA1007121//INOSITOL POLYPHOSPHATE 1-PHOSPHATASE (EC 3.1.3.57) (IPP). //5.4e-07:90:28//HOMO SAPIENS (HUMAN). //P49441
 F-HEMBA1007129//HIRUSTASIN. //0.88:37:32//HIRUDO MEDICINALIS (MEDICINAL LEECH). //P80302
 F-HEMBA1007147//HYPOTHETICAL 12.0 KD PROTEIN IN DST1-HEM2 INTERGENIC REGION. //0.92:23:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53182
 F-HEMBA1007149//BACTERIOCIN MICROCIN B17 PRECURSOR (MCB17). //0.0078:17:70//ESCHERICHIA COLI. //P05834
 F-HEMBA1007151//WDM1 PROTEIN PRECURSOR. //0.25:45:37//MUS MUSCULUS (MOUSE). //Q62477
 F-HEMBA1007174//HYPOTHETICAL 45.1 KD PROTEIN IN RP55-ZMS1 INTERGENIC REGION. //6.9e-18:97:47//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P47160
 F-HEMBA1007178//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!/9.8e-06:38:65//HOMO SAPIENS (HUMAN). //P39195
 F-HEMBA1007194//GLUCOSE-6-PHOSPHATE 1-DEHYDROGENASE, CHLOROPLAST ISOFORM PRECURSOR (EC 1.1.1.49) (G6PD). //1.0:80:32//NICOTIANA TABACUM (COMMON TOBACCO). //Q43793
 F-HEMBA1007203//PROTEIN A22. //1.0:115:26//VARIOLA VIRUS. //P33845
 F-HEMBA1007206
 F-HEMBA1007224//HYPOTHETICAL 35.7 KD PROTEIN C41C 4.6 IN CHROMOSOME 11. //2.4e-05:92:30//CAENORHABDITIS ELEGANS. //Q09275
 F-HEMBA1007243//HYPOXANTHINE-GUANINE PHOSPHORIBOSYLTRANSFERASE (EC 2.4.2.8) (HGPRT) (HGPRTASE) (HPRT B). //3.1e-74:205:67//MUS MUSCULUS (MOUSE). //P00493
 F-HEMBA1007251//VITELLINE MEMBRANE PROTEIN VM26AB PRECURSOR (PROTEIN TU-4) (PROTEIN SV23). //0.52:108:30//DROSOPHILA MELANOGASTER (FRUIT FLY). //P13238
 F-HEMBA1007256
 10 F-HEMBA1007267//CALICIN (FRAGMENT). //0.060:88:31//HOMO SAPIENS (HUMAN). //Q13939
 F-HEMBA1007273//HYPOTHETICAL 8.1 KD PROTEIN (ORF6 5). //0.95:40:37//GUILLARDIA THETA (CRYPTOMONAS PH1). //078421
 F-HEMBA1007279//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!/4.6e-24:98:64//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1007281
 F-HEMBA1007288//HYPOTHETICAL 13.5 KD PROTEIN IN ZMS1-MNS1 INTERGENIC REGION. //0.88:11:54//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P47162
 F-HEMBA1007300//CGMP-SPECIFIC 3',5'-CYCLIC PHOSPHODIESTERASE (EC 3.1.4.17) (CGB-PDE). //2.7e-43:220:41//BOS TAURUS (BOVINE). //Q28156
 F-HEMBA1007301//PROCOLLAGEN ALPHA 1(I11) CHAIN PRECURSOR. //3.3e-22:115:33//HOMO SAPIENS (HUMAN). //P02461
 F-HEMBA1007319
 F-HEMBA1007320//HYPOTHETICAL 28.0 KD PROTEIN IN GLOB-RNHA INTERGENIC REGION. //1.0:48:37//ESCHERICHIA COLI. //P75672
 F-HEMBA1007322//THREONINE DEHYDRATASE OPERON ACTIVATOR PROTEIN. //1.0:59:33//ESCHERICHIA COLI. //P11866
 F-HEMBA1007327
 F-HEMBA1007341//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!/9.1e-12:37:62//HOMO SAPIENS (HUMAN). //P39188
 F-HEMBA1007342//PROBABLE E5 PROTEIN. //0.89:96:29//PYGMY CHIMPANZEE PAPILOMAVIRUS TYPE 1. //Q02268
 F-HEMBA1007347//INSULIN-LIKE GROWTH FACTOR BINDING PROTEIN 2 PRECURSOR (IGFBP-2) (IBP-2) (IGF-BINDING PROTEIN 2). //0.92:62:43//OVIS ARIES (SHEEP). //Q29400
 F-HEMBA1000005//WEAK NEUROTOXIN 5. //0.98:30:33//NAJAJA NAJA (INDIAN COBRA). //P29179
 F-HEMBA1000008//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!/2.7e-35:73:84//HOMO SAPIENS (HUMAN). //P39195
 F-HEMBA1000018//HYPOTHETICAL BHLF1 PROTEIN. //0.39:

90:37//EPSTEIN-BARR VIRUS (STRAIN B95-8) (HUMAN HE
RPESVIRUS 4).//P03181
F-HEMBB1000024//VIRE LOCUS 9 KD VIRULENCE PROTEIN.
//0.66:36:41//AGROBACTERIUM TUMEFACIENS.//P08061
F-HEMBB1000025//MUSCARINIC TOXIN ALPHA (MT-ALPHA).
//0.46:32:40//DENDROASPIS POLYLEPIS POLYLEPIS (BLA
CK MAMBA).//P80494
F-HEMBB1000030//SUPPRESSOR PROTEIN SRP40.//6.7e-0
7:50:52//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST)./
/P32583
F-HEMBB1000036//HYPOTHETICAL 43.2 KD PROTEIN C34E1
0.1 IN CHROMOSOME III.//2.5e-07:120:29//CAENORHABDI
TIS ELEGANS.//P46576
F-HEMBB1000037//HYPOTHETICAL 59.9 KD PROTEIN IN SG
A1-KTR7 INTERGENIC REGION.//1.7e-05:71:29//SACCHAR
OMYCES CEREVISIAE (BAKER'S YEAST).//P40492
F-HEMBB1000039//VERY HYPOTHETICAL 11.9 KD PROTEIN
C4H3.12C IN CHROMOSOME I.//1.0:61:21//SCHIZOSACCHAR
OMYCES POMBE (FISSION YEAST).//Q10219
F-HEMBB1000044
F-HEMBB1000048//HYPOTHETICAL 15.7 KD PROTEIN IN ID
H-DEOR INTERGENIC REGION.//1.0:63:31//BACILLUS SUB
TILIS.//P54942
F-HEMBB1000050//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/9.0e-14:34:79//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBB1000054//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/5.9e-31:45:73//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBB1000055//MUSCARINIC TOXIN ALPHA (MT-ALPHA).
//1.0:14:57//DENDROASPIS POLYLEPIS POLYLEPIS (BLAC
K MAMBA).//P80494
F-HEMBB1000059//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/1.0e-21:82:59//HOMO SAPIENS (HUMAN).//P391
95
F-HEMBB1000083//CHROMOGANIN A PRECURSOR (CGA) [CO
NTAINS: PANCREASTATIN; BETA-GRANIN; WE-14].//0.87:1
72:28//RATTUS NORVEGICUS (RAT).//P10354
F-HEMBB1000089//HYPOTHETICAL 9.5 KD PROTEIN IN SPE
A-METK INTERGENIC REGION (F83).//1.0:42:33//ESCHER
ICHIA COLI.//P46879
F-HEMBB1000099//!!!! ALU SUBFAMILY SB WARNING ENTR
Y !!!!!/7.7e-08:31:87//HOMO SAPIENS (HUMAN).//P391
89
F-HEMBB1000103//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//1.4e-38:136:58//HOMO SAPIENS (HUMAN).//P08547
F-HEMBB1000113//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/3.9e-13:57:64//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBB1000119//MAF PROTEIN.//3.6e-32:195:43//BACI

LLUS SUBTILIS.//Q02169
F-HEMBB1000136//HYPOTHETICAL 12.7 KD PROTEIN IN PC
S60-ABD1 INTERGENIC REGION.//0.65:71:32//SACCHAROM
YCES CEREVISIAE (BAKER'S YEAST).//P38327
F-HEMBB1000141//B-CELL GROWTH FACTOR PRECURSOR (BC
GF-12 KD).//0.00014:34:64//HOMO SAPIENS (HUMAN).//
P20931
F-HEMBB1000144//!!!! ALU SUBFAMILY SB2 WARNING ENT
RY !!!!!/2.0e-26:81:69//HOMO SAPIENS (HUMAN).//P39
10 191
F-HEMBB1000173//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/9.2e-29:91:71//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBB1000175//ANTIMICROBIAL PEPTIDE ENAP-1 (FRAG
MENT).//0.97:41:36//EQUUS CABALLUS (HORSE).//P8093
0
F-HEMBB1000198//HYPOTHETICAL 7.7 KD PROTEIN YCF33
(ORF67).//0.91:21:52//PORPHYRA PURPUREA.//P51329
F-HEMBB1000215//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!/3.4e-08:39:76//HOMO SAPIENS (HUMAN).//P391
92
F-HEMBB1000217//DNA DAMAGE TOLERANCE PROTEIN RHC31
(RAD31 HOMOLOG).//2.9e-32:174:40//SACCHAROMYCES C
EREVISIAE (BAKER'S YEAST).//Q06624
F-HEMBB1000218//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//0.73:31:38//MICROTUS PENNSYLVANICUS (ME
ADOW VOLE).//P24949
F-HEMBB1000226//HYPOTHETICAL 37.0 KD PROTEIN B049
5.8 IN CHROMOSOME II.//6.5e-26:191:34//CAENORHABDI
TIS ELEGANS.//Q09217
F-HEMBB1000240
F-HEMBB1000244//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/1.9e-05:44:61//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBB1000250
F-HEMBB1000258
F-HEMBB1000264//CUTICLE COLLAGEN SQT-1.//0.15:89:3
3//CAENORHABDITIS ELEGANS.//P12114
F-HEMBB1000266//TRANSLATION INITIATION FACTOR IF-
2.//2.7e-06:167:22//SACCHAROMYCES CEREVISIAE (BAKE
R'S YEAST).//P39730
F-HEMBB1000272//CYTOCHROME C OXIDASE POLYPEPTIDE V
IB (EC 1.9.3.1) (AED).//0.75:30:43//BOS TAURUS (BO
VINE).//P00429
F-HEMBB1000274//CORNIFIN (SMALL PROLINE-RICH PROTE
IN I) (SPR-I) (SMALL PROLINE-RICH SQUAMOUS CELL MA
RKER) (SPRP).//1.0:38:36//SUS SCROFA (PIG).//P3532
3
F-HEMBB1000284//CALTRIN (CALCIUM TRANSPORT INHIBIT
OR).//1.0:56:30//MUS MUSCULUS (MOUSE).//Q09098

- F-HEMBB1000307
 F-HEMBB1000312
 F-HEMBB1000317//THROMBOSPONDIN 1 PRECURSOR.//3.2e-32:135:43//HOMO SAPIENS (HUMAN).//P07996
 F-HEMBB1000318//PUTATIVE SMALL MEMBRANE PROTEIN (NONSTRUCTURAL PROTEIN NS3) (NONSTRUCTURAL 9.5 KD PROTEIN).//0.41:51:31//HUMAN CORONAVIRUS (STRAIN OC43).//Q04854
 F-HEMBB1000335//ZINC FINGER PROTEIN 13 (ZFP-13) (KROX-8 PROTEIN) (FRAGMENT).//0.82:33:45//MUS MUSCULUS (MOUSE).//P10754
 F-HEMBB1000336//ALDEHYDE OXIDASE (EC 1.2.3.1) (FRAGMENTS).//0.80:44:40//ORYZOLAGUS CUNICULUS (RABBIT).//P80456
 F-HEMBB1000337//SPLICING FACTOR, ARGININE/SERINE-RICH 4 (PRE-MRNA SPLICING FACTOR SRP75).//0.94:118:22//HOMO SAPIENS (HUMAN).//Q08170
 F-HEMBB1000338//MALE SPECIFIC SPERM PROTEIN MST84D A.//0.042:33:39//DROSOPHILA MELANOGASTER (FRUIT FL Y).//Q01642
 F-HEMBB1000339//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/2.2e-14:54:55//HOMO SAPIENS (HUMAN).//P39188
 F-HEMBB1000341//GENE 74 PROTEIN (GP74).//1.0:39:33//MYCOBACTERIOPHAGE L5.//Q05289
 F-HEMBB1000343
 F-HEMBB1000354//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/1.1e-15:83:56//HOMO SAPIENS (HUMAN).//P39188
 F-HEMBB1000369//PROTEIN Q300.//0.99:27:40//MUS MUSCULUS (MOUSE).//Q02722
 F-HEMBB1000374//!!!! ALU SUBFAMILY SB WARNING ENTRY
 Y !!!!!/4.7e-34:56:78//HOMO SAPIENS (HUMAN).//P39189
 F-HEMBB1000376
 F-HEMBB1000391//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//0.0013:79:35//BOS TAURUS (BOVINE).//P25508
 F-HEMBB1000399//CHECKPOINT PROTEIN RAD17.//2.8e-15:187:31//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//P50531
 F-HEMBB1000402//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 5 (EC 1.6.5.3) (FRAGMENT).//0.027:60:38//LEISHMANIA TARENTOLAE (SAUROLEISHMANIA TARENTOLAE).//P15583
 F-HEMBB1000404//CYANELLE 50S RIBOSOMAL PROTEIN L28.//0.94:29:27//CYANOPHORA PARADOXA.//P48129
 F-HEMBB1000420//SPLICEOSOME ASSOCIATED PROTEIN 49 (SAP 49) (SF3B53).//0.023:97:35//HOMO SAPIENS (HUMAN).//Q15427
 F-HEMBB1000434//!!!! ALU SUBFAMILY SQ WARNING ENTRY
- Y !!!!!/4.8e-20:111:54//HOMO SAPIENS (HUMAN).//P39194
 F-HEMBB1000438//HYPOTHETICAL 7.9 KD PROTEIN IN GP55-NRDG INTERGENIC REGION.//0.93:24:50//BACTERIOPHAGE T4.//P07076
 F-HEMBB1000441//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/4.4e-23:85:70//HOMO SAPIENS (HUMAN).//P39188
 F-HEMBB1000449//!!!! ALU SUBFAMILY SX WARNING ENTRY
 Y !!!!!/0.88:27:51//HOMO SAPIENS (HUMAN).//P39195
 F-HEMBB1000455
 F-HEMBB1000472
 F-HEMBB1000480//PROTEIN STBC.//1.0:52:30//ESCHERICHIA COLI.//P11905
 F-HEMBB1000487//SHORT NEUROTOXIN 1 (NEUROTOXIN ALPHA) (NEUROTOXIN II).//0.93:29:34//NAJA OXIANA (CENTRAL ASIAN COBRA) (OXUS COBRA).//P01427
 F-HEMBB1000490//!!!! ALU SUBFAMILY SX WARNING ENTRY
 Y !!!!!/2.3e-16:50:80//HOMO SAPIENS (HUMAN).//P39195
 F-HEMBB1000491
 F-HEMBB1000493//3A PROTEIN.//1.0:51:35//AVIAN INFECTIOUS BRONCHITIS VIRUS (STRAIN BEAUDETTE) (IBV).//P30237
 F-HEMBB1000510//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//9.7e-27:132:45//HOMO SAPIENS (HUMAN).//P08547
 F-HEMBB1000518//CYTOCHROME C OXIDASE POLYPEPTIDE I II (EC 1.9.3.1).//0.021:47:40//LEISHMANIA TARENTOLAE (SAUROLEISHMANIA TARENTOLAE).//P14546
 F-HEMBB1000523
 F-HEMBB1000530//COLLAGEN ALPHA 1(XIV) CHAIN PRECURSOR (UNDULIN).//9.8e-14:43:83//GALLUS GALLUS (CHICKEN).//P32018
 F-HEMBB1000550//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 5 (EC 1.6.5.3).//0.19:97:30//TRYPANOSOMA BRUCEI
 BRUCEI.//P04540
 F-HEMBB1000554//MATERNAL B9.10 PROTEIN (P30 B9.10).//0.94:82:25//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P40744
 F-HEMBB1000556//TRANSCRIPTION INITIATION FACTOR TFIID 135 KD SUBUNIT (TAFII-135) (TAFII135) (TAFII-130) (TAFII130).//0.043:201:29//HOMO SAPIENS (HUMAN).//000268
 F-HEMBB1000564//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (A6L).//1.0:52:34//METRIDIVM SENILE (BROWN SEA ANEMONE) (FRILLED SEA ANEMONE).//047493
 F-HEMBB1000573//!!!! ALU SUBFAMILY SB2 WARNING ENTRY
 Y !!!!!/2.3e-10:52:73//HOMO SAPIENS (HUMAN).//P39191
 F-HEMBB1000575//!!!! ALU SUBFAMILY SC WARNING ENTRY

2917

Y !!!!!/1.8e-26:76:76//HOMO SAPIENS (HUMAN).//P391
92
F-HEMBB1000586//NADH-UBIQUINONE OXIDOREDUCTASE MLR
Q SUBUNIT (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-MLR
Q) (CI-MLRQ).//0.74:23:52//HOMO SAPIENS (HUMAN).//
000483
F-HEMBB1000589//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/2.9e-25:61:75//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBB1000591//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//1.0:34:35//PETROMYZON MARINUS (SEA LAMP
REY).//Q35537
F-HEMBB1000592//SMALL PROLINE-RICH PROTEIN 2-1.//
0.0016:49:42//HOMO SAPIENS (HUMAN).//P35326
F-HEMBB1000593//COLLAGEN ALPHA 1(III) CHAIN (FRAGM
ENTS).//0.0070:189:32//GALLUS GALLUS (CHICKEN).//P
12105
F-HEMBB1000598//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//5.7e-10:110:41//NYCTICEBUS COUCANG (SLOW LORI
S).//P08548
F-HEMBB1000623//HYPOTHETICAL 54.9 KD PROTEIN C02F
5.7 IN CHROMOSOME III.//0.0022:98:28//CAENORHABDIT
IS ELEGANS.//P34284
F-HEMBB1000630
F-HEMBB1000631//ALPHA-2C-1 ADRENERGIC RECEPTOR (AL
PHA-2C-1 ADRENOCEPTOR) (SUBTYPE C4).//8.8e-06:59:40
//HOMO SAPIENS (HUMAN).//P18825
F-HEMBB1000632//GUANINE NUCLEOTIDE RELEASING PROTE
IN (GNRP).//7.3e-13:173:28//MUS MUSCULUS (MOUSE).//
P27671
F-HEMBB1000637//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/4.6e-41:94:82//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBB1000638//INVOLUCRIN.//1.9e-06:144:29//HOMO
SAPIENS (HUMAN).//P07476
F-HEMBB1000643//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/8.3e-30:77:76//HOMO SAPIENS (HUMAN).//P391
95
F-HEMBB1000649//!!!! ALU SUBFAMILY SB WARNING ENTR
Y !!!!!/2.5e-37:58:81//HOMO SAPIENS (HUMAN).//P391
89
F-HEMBB1000652//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/2.2e-37:61:77//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBB1000665//HYPOTHETICAL PROTEIN BBD24.//0.83:
38:36//BORRELIA BURGDORFERI (LYME DISEASE SPIROCHE
TE).//P70845
F-HEMBB1000671//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//6.8e-51:74:71//HOMO SAPIENS (HUMAN).//P08547
F-HEMBB1000673//HEAT-STABLE ENTEROTOXIN A3/A4 PREC

2918

URSOR (STA3/STA4) (ST-IB) (ST-H).//0.012:37:37//ES
CHERICHIA COLI.//P07965
F-HEMBB1000684//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/3.1e-21:66:72//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBB1000693//HUNTINGTIN ASSOCIATED PROTEIN 1 (H
AP1).//5.2e-26:121:49//RATTUS NORVEGICUS (RAT).//P
54256
F-HEMBB1000705
F-HEMBB1000706
F-HEMBB1000709//HYPOTHETICAL 5.8 KD PROTEIN.//1.0:
29:44//CLOVER YELLOW MOSAIC VIRUS (CYMV).//P16485
F-HEMBB1000725//RAS-RELATED PROTEIN RAB-8B.//7.4e-
105:205:98//RATTUS NORVEGICUS (RAT).//P70550
F-HEMBB1000726//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/1.4e-25:85:70//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBB1000738//50S RIBOSOMAL PROTEIN L33.//1.0:4
1:31//THERMUS AQUATICUS (SUBSP. THERMOPHILUS).//P35
20 871
F-HEMBB1000749//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/1.3e-29:42:85//HOMO SAPIENS (HUMAN).//P391
94
F-HEMBB1000763//NIFU PROTEIN.//0.089:63:36//FRANKI
A ALNI.//P46045
F-HEMBB1000770//CALTRIN-LIKE PROTEIN II.//0.98:13:
69//CAVIA PORCELLUS (GUINEA PIG).//P22075
F-HEMBB1000774//HIGH MOBILITY GROUP PROTEIN HMG-Y.
//0.029:53:32//MUS MUSCULUS (MOUSE).//P17095
30 F-HEMBB1000781//MAPK/ERK KINASE KINASE 2 (EC 2.7.
1.-) (MEK KINASE 2) (MEKK 2).//3.5e-75:144:98//MUS
MUSCULUS (MOUSE).//Q61083
F-HEMBB1000789//PUTATIVE 90.2 KD ZINC FINGER PROTE
IN IN CCA1-ADK2 INTERGENIC REGION.//2.6e-49:232:43
//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P3995
6
F-HEMBB1000790//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/2.9e-16:93:51//HOMO SAPIENS (HUMAN).//P3918
8
F-HEMBB1000794
F-HEMBB1000807//MUSCARINIC ACETYLCHOLINE RECEPTOR
M3.//0.54:111:27//GALLUS GALLUS (CHICKEN).//P49578
F-HEMBB1000810
F-HEMBB1000821
F-HEMBB1000822//HYPOTHETICAL 10 KD PROTEIN (ORF
6).//0.10:50:34//NARCISSUS MOSAIC VIRUS (NMV).//P1
5099
F-HEMBB1000826//B-CELL GROWTH FACTOR PRECURSOR (BC
GF-12 KD).//0.00025:73:39//HOMO SAPIENS (HUMAN).//
P20931

F-HEMBB1000827//HYPOTHETICAL 7.4 KD PROTEIN.//0.8
9:23:52//THERMOPROTEUSTENAX VIRUS 1 (STRAIN KRA1)
(TTV1).//P19302
F-HEMBB1000831//MALE SPECIFIC SPERM PROTEIN MST87
F.//0.98:35:40//DROSOPHILA MELANOGASTER (FRUIT FL
Y).//P08175
F-HEMBB1000835//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//7.8e-31:96:46//HOMO SAPIENS (HUMAN).//P08547
F-HEMBB1000840//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//0.00012:102:36//NYCTICEBUS COUCANG (SLOW LORI
S).//P08548
F-HEMBB1000848//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//7.3e-97:239:70//HOMO SAPIENS (HUMAN).//P08547
F-HEMBB1000852
F-HEMBB1000870
F-HEMBB1000876//METALLOTHIONEIN (MT).//0.99:14:64/
/PERCA FLUVIATILIS (PERCH).//P52725
F-HEMBB1000883//HYPOTHETICAL 7.8 KD PROTEIN (ORF6
2).//0.34:60:33//GUILLARDIA THETA (CRYPTOMONAS PH
I).//078459
F-HEMBB1000887//HISTIDINE-RICH, METAL BINDING POLY
PEPTIDE.//1.0:26:42//HELICOBACTER PYLORI (CAMPYLO
BACTER PYLORI).//Q48251
F-HEMBB1000888
F-HEMBB1000890
F-HEMBB1000893
F-HEMBB1000908//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//0.0074:45:51//HOMO SAPIENS (HUMAN).//P39188
F-HEMBB1000910//PROBABLE E5 PROTEIN.//1.0:49:36//H
UMAN PAPILLOMAVIRUS TYPE 58.//P26552
F-HEMBB1000913//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!//0.29:56:46//HOMO SAPIENS (HUMAN).//P39195
F-HEMBB1000915//CYTOCHROME B (EC 1.10.2.2).//2.5e-
24:62:90//HOMO SAPIENS (HUMAN).//P00156
F-HEMBB1000917//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!//5.9e-26:53:66//HOMO SAPIENS (HUMAN).//P391
93
F-HEMBB1000927//NEURONAL CALCIUM SENSOR 1 (NCS-1)
(FREQUENIN).//3.9e-44:182:45//XENOPUS LAEVIS (AFRI
CAN CLAWED FROG).//Q91614
F-HEMBB1000947//SMALL PROLINE-RICH PROTEIN 2-1.//
0.24:69:27//HOMO SAPIENS (HUMAN).//P35326
F-HEMBB1000959//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!//3.0e-31:89:68//HOMO SAPIENS (HUMAN).//P391
95
F-HEMBB1000973//CONNECTIVE TISSUE GROWTH FACTOR PR
ECURSOR.//0.96:66:36//BOS TAURUS (BOVINE).//018739
F-HEMBB1000975//HISTIDINE-RICH GLYCOPROTEIN PRECUR
SOR (HISTIDINE-PROLINERICH GLYCOPROTEIN) (HPRG).//
0.00042:77:41//HOMO SAPIENS (HUMAN).//P04196

F-HEMBB1000981
F-HEMBB1000985//MIPP PROTEIN (MURINE IAP-PROMOTED
PLACENTA-EXPRESSED PROTEIN).//1.0e-18:178:30//MUS
MUSCULUS (MOUSE).//P28575
F-HEMBB1000991
F-HEMBB1000996//HYPOTHETICAL 10.1 KD PROTEIN IN RH
SD-CCL INTERGENIC REGION (ORFD3).//0.58:34:35//ESC
HERICHA COLI.//P33669
F-HEMBB1001004//PROBABLE E4 PROTEIN.//0.24:110:35/
/HUMAN PAPILLOMAVIRUS TYPE 5B.//P26550
F-HEMBB1001008
F-HEMBB1001011//ZINC FINGER PROTEIN 7 (ZINC FINGER
PROTEIN KOX4) (ZINC FINGER PROTEIN HF.16).//3.2e-
17:104:47//HOMO SAPIENS (HUMAN).//P17097
F-HEMBB1001014//EOTAXIN PRECURSOR (EOSINOPHIL CHEM
OTACTIC PROTEIN).//1.0:58:39//RATTUS NORVEGICUS (R
AT).//P97545
F-HEMBB1001020//!!!! ALU SUBFAMILY SB WARNING ENTR
Y !!!!!//1.4e-07:36:75//HOMO SAPIENS (HUMAN).//P391
89
F-HEMBB1001024
F-HEMBB1001037//FERREDOXIN.//1.0:52:25//MOORELLA T
HERMOACETICA (CLOSTRIDIUM THERMOACETICUM).//P00203
F-HEMBB1001047
F-HEMBB1001051//PROTEIN FAN (FACTOR ASSOCIATED WIT
H N-SMASE ACTIVATION).//3.4e-21:50:100//HOMO SAPI
ENS (HUMAN).//Q92636
F-HEMBB1001056//HYPOTHETICAL 29.3 KD PROTEIN (ORF9
2).//0.0099:115:35//ORGYIA PSEUDOTSUGATA MULTICAPS
ID POLYHEDROSIS VIRUS (OPMNPV).//010341
F-HEMBB1001058//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!//1.1e-33:95:76//HOMO SAPIENS (HUMAN).//P391
92
F-HEMBB1001060//HYPOTHETICAL 8.2 KD PROTEIN ZC21.7
IN CHROMOSOME III.//1.0:38:36//CAENORHABDITIS ELE
GANS.//P34591
F-HEMBB1001063
F-HEMBB1001068
F-HEMBB1001096//NOXIUSTOXIN (NTX) (TOXIN II.11).//
0.99:36:38//CENTRUROIDES NOXIUS (MEXICAN SCORPIO
N).//P08815
F-HEMBB1001102//HYPOTHETICAL 72.5 KD PROTEIN C2F7.
10 IN CHROMOSOME I.//1.1e-27:115:36//SCHIZOSACCHAR
OMYCES POMBE (FISSION YEAST).//Q09701
F-HEMBB1001105//CLASS II HISTOCOMPATIBILITY ANTIGE
N, M ALPHA CHAIN PRECURSOR.//0.80:70:40//HOMO SAPI
ENS (HUMAN).//P28067
F-HEMBB1001112//PROTEIN TRANSPORT PROTEIN SEC61 AL
PHA SUBUNIT.//1.1e-126:287:85//RATTUS NORVEGICUS
(RAT).//P38378

2921

F-HEM BB1001114//HYPOTHETICAL 9.6 KD PROTEIN (ORF 2). //0.84:62:27//BACTERIOPHAGE L2. //P42537
 F-HEM BB1001117
 F-HEM BB1001119//COLLAGEN ALPHA 1(XII) CHAIN PRECURSOR. //1.6e-21:50:98//HOMO SAPIENS (HUMAN). //Q99715
 F-HEM BB1001126//HYPOTHETICAL 55.9 KD PROTEIN EED 8.6 IN CHROMOSOME II. //1.7e-50:184:53//CAENORHABDITIS ELEGANS. //Q09296
 F-HEM BB1001133//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!//1.4e-09:53:62//HOMO SAPIENS (HUMAN). //P391 10 92
 F-HEM BB1001137//DNA-DIRECTED RNA POLYMERASE II LARGE SUBUNIT (EC 2.7.7.6) (RPB1) (FRAGMENT). //2.0e-05:206:27//CRICETULUS GRISEUS (CHINESE HAMSTER). //P11414
 F-HEM BB1001142//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//4.1e-05:46:56//HOMO SAPIENS (HUMAN). //P391 93
 F-HEM BB1001151//HYPOTHETICAL 33.5 KD PROTEIN C1D4. 02C IN CHROMOSOME I. //2.3e-23:109:44//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10149
 F-HEM BB1001153//PROCOLLAGEN ALPHA 2(IV) CHAIN PRECURSOR. //0.75:76:34//ASCARIS SUUM (PIC ROUNDWORM) (ASCARIS LUMBRICOIDES). //P27393
 F-HEM BB1001169//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//1.4e-16:71:59//HOMO SAPIENS (HUMAN). //P391 95
 F-HEM BB1001175//ANKYRIN. //3.2e-12:169:31//MUS MUSCULUS (MOUSE). //Q02357
 F-HEM BB1001177//PERIODIC TRYPTOPHAN PROTEIN 2 HOMOLOG. //9.4e-07:148:27//HOMO SAPIENS (HUMAN). //Q1526 9
 F-HEM BB1001182//HYPOTHETICAL 36.0 KD PROTEIN. //1.3e-09:110:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P54858
 F-HEM BB1001199
 F-HEM BB1001208//HYPOTHETICAL PROTEIN LAMBDA-SP5. //0.053:23:47//MUS MUSCULUS (MOUSE). //P15974
 F-HEM BB1001209
 F-HEM BB1001210//HYPOTHETICAL PROTEIN LAMBDA-SP5. // 40 0.14:40:37//MUS MUSCULUS (MOUSE). //P15974
 F-HEM BB1001218//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//3.4e-19:49:67//HOMO SAPIENS (HUMAN). //P391 94
 F-HEM BB1001221//CYTOCHROME C OXIDASE POLYPEPTIDE V IIA-LIVER PRECURSOR (EC 1.9.3.1). //0.11:44:38//HOMO SAPIENS (HUMAN). //P14406
 F-HEM BB1001234//65 KD YES-ASSOCIATED PROTEIN (YAP65). //2.0e-45:192:53//MUS MUSCULUS (MOUSE). //P46938
 F-HEM BB1001242//HYPOTHETICAL 143.3 KD TRP-ASP REPE 50

2922

ATS CONTAINING PROTEIN C12G12.13C IN CHROMOSOME I. //5.5e-37:226:41//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09876
 F-HEM BB1001249//OXALOACETATE DECARBOXYLASE GAMMA CHAIN (EC 4.1.1.3). //1.0:23:43//KLEBSIELLA PNEUMONIAE. //P13155
 F-HEM BB1001253//METALLOTHIONEIN-1H (MT-1H) (METALLOTHIONEIN-0) (MT-0). //0.14:16:43//HOMO SAPIENS (HUMAN). //P80294
 F-HEM BB1001254//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//1.4e-12:40:75//HOMO SAPIENS (HUMAN). //P391 95
 F-HEM BB1001267//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//1.0e-12:33:78//HOMO SAPIENS (HUMAN). //P391 93
 F-HEM BB1001271//HYPOTHETICAL 25.1 KD PROTEIN B030 2.5 IN CHROMOSOME X. //1.0:58:37//CAENORHABDITIS ELEGANS. //Q10928
 F-HEM BB1001282//ANKYRIN HOMOLOG PRECURSOR. //9.5e-13:206:31//CHROMATIUM VINOSUM. //Q06527
 F-HEM BB1001288//COPPER HOMEOSTASIS PROTEIN CUTC. //4.6e-42:163:51//ESCHERICHIA COLI. //P46719
 F-HEM BB1001289//HYPOTHETICAL PROTEIN ORF-1137. //1.0e-05:106:26//MUS MUSCULUS (MOUSE). //P11260
 F-HEM BB1001294//GTP-BINDING PROTEIN TC10. //1.3e-34:58:94//HOMO SAPIENS (HUMAN). //P17081
 F-HEM BB1001302//HOMEBOX PROTEIN CDX-2 (CAUDAL-TYPE HOMEBOX PROTEIN 2) (CDX-3). //0.24:49:46//HOMO SAPIENS (HUMAN). //Q99626
 F-HEM BB1001304//GLYCINE-RICH CELL WALL STRUCTURAL PROTEIN (CLONE W10-1) (FRAGMENT). //1.0:17:70//LYCOPERSICON ESCULENTUM (TOMATO). //Q01157
 F-HEM BB1001314//SALIVARY GLUE PROTEIN SGS-3 PRECURSOR. //0.21:104:27//DROSOPHILA ERECTA (FRUIT FLY). //P13730
 F-HEM BB1001315//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//1.3e-24:53:71//HOMO SAPIENS (HUMAN). //P391 95
 F-HEM BB1001317//HYPOTHETICAL 85.7 KD PROTEIN C13G 6.03 IN CHROMOSOME I. //0.24:90:31//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09782
 F-HEM BB1001326//HYPOTHETICAL PROTEIN LAMBDA-SP5. //0.36:26:50//MUS MUSCULUS (MOUSE). //P15974
 F-HEM BB1001331//HYPOTHETICAL BHLF1 PROTEIN. //1.0:127:33//EPSTEIN-BARR VIRUS (STRAIN B95-8) (HUMAN HERPESVIRUS 4). //P03181
 F-HEM BB1001335//ESCARGOT/SLAIL PROTEIN HOMOLOG (FRAGMENT). //0.85:44:29//SCIARA COPROPHILA (FUNGUS GNAT). //Q01799
 F-HEM BB1001337//!!!! ALU SUBFAMILY SQ WARNING ENTRY

2923

Y !!!!!/4.2e-20:62:62//HOMO SAPIENS (HUMAN).//P391
94
F-HEM BB1001339//HYPOTHETICAL 17.3 KD PROTEIN CY1A1
1.16C.//8.2e-07:123:34//MYCOBACTERIUM TUBERCULOSI
S.//Q50606
F-HEM BB1001346//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//4.2e-14:60:45//HOMO SAPIENS (HUMAN).//P08547
F-HEM BB1001348//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/5.6e-14:61:62//HOMO SAPIENS (HUMAN).//P3918
8
F-HEM BB1001356
F-HEM BB1001364
F-HEM BB1001366//HISTIDINE-RICH PROTEIN.//0.87:26:4
2//PLASMODIUM FALCIPARUM (ISOLATE FCM17 / SENECA
L).//P14586
F-HEM BB1001367//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!/8.6e-40:146:61//HOMO SAPIENS (HUMAN).//P39
192
F-HEM BB1001369
F-HEM BB1001380//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/1.9e-25:49:83//HOMO SAPIENS (HUMAN).//P391
93
F-HEM BB1001384//BH3 INTERACTING DOMAIN DEATH AGONI
ST (BID).//0.80:95:29//MUS MUSCULUS (MOUSE).//P704
44
F-HEM BB1001387//PEA2 PROTEIN (PPF2 PROTEIN).//0.02
2:117:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAS
T).//P40091
F-HEM BB1001394//ALPHA-ADAPTIN A (CLATHRIN ASSEMBLY
PROTEIN COMPLEX 2 ALPHA-A LARGE CHAIN) (100 KD CO
ATED VESICLE PROTEIN A) (PLASMA MEMBRANE ADAPTOR H
A2/AP2 ADAPTIN ALPHA A SUBUNIT).//0.38:85:31//MUS
MUSCULUS (MOUSE).//P17426
F-HEM BB1001410
F-HEM BB1001424//PHOTOSYSTEM II 4 KD REACTION CENTR
E PROTEIN PRECURSOR.//0.99:37:21//ORYZA SATIVA (RI
CE).//P12162
F-HEM BB1001426//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/0.0035:40:60//HOMO SAPIENS (HUMAN).//P3919
5
F-HEM BB1001429//CYTOSOL AMINOPEPTIDASE (EC 3.4.11.
1) (LEUCINE AMINOPEPTIDASE) (LAP) (LEUCYL AMINOPEP
TIDASE) (PROLINE AMINOPEPTIDASE) (EC 3.4.11.5) (PR
OLYL AMINOPEPTIDASE).//1.1e-99:219:86//BOS TAURUS
(BOVINE).//P00727
F-HEM BB1001436//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/3.4e-30:57:78//HOMO SAPIENS (HUMAN).//P391
95
F-HEM BB1001443//[PYRUVATE DEHYDROGENASE (LIPOAMID
E)]-PHOSPHATASE PRECURSOR (PDP) (EC 3.1.3.43) (PYR

2924

UVATE DEHYDROGENASE PHOSPHATASE, CATALYTIC SUBUNI
T) (PDPC).//2.5e-79:155:97//BOS TAURUS (BOVINE).//
P35816
F-HEM BB1001449
F-HEM BB1001454//SALIVARY PROLINE-RICH PROTEIN PO
(ALLELE M) [CONTAINS: PEPTIDE P-D] (FRAGMENT).//1.
1e-05:196:31//HOMO SAPIENS (HUMAN).//P10161
F-HEM BB1001458//24 KD ANTIGEN (FRAGMENT).//0.94:1
8:50//PLASMODIUM CHABAUDI.//P14592
10 F-HEM BB1001463
F-HEM BB1001464//PPF2L ANTIGEN (FRAGMENT).//1.0:45:
28//PLASMODIUM FALCIPARUM (ISOLATE PALO ALTO / UGA
NDA).//P07765
F-HEM BB1001482//GASTRULA ZINC FINGER PROTEIN XLCCF
16.1 (FRAGMENT).//4.2e-10:37:43//XENOPUS LAEVIS (A
FRICAN CLAWED FROG).//P18712
F-HEM BB1001500
F-HEM BB1001521//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/2.4e-39:59:72//HOMO SAPIENS (HUMAN).//P3918
8
F-HEM BB1001527//HOMEBOX PROTEIN HOX-B5 (XLHBOX-4)
(XHOX-1B) (FRAGMENT).//0.21:131:25//XENOPUS LAEVI
S (AFRICAN CLAWED FROG).//P09019
F-HEM BB1001531//GENE 32 PROTEIN (GP32).//0.88:95:3
0//MYCOBACTERIOPHAGE L5.//Q05241
F-HEM BB1001535//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (AGL).//1.0:31:38//LUMBRICUS TERRESTRIS (COMMON
EARTHWORM).//Q34942
F-HEM BB1001536
30 F-HEM BB1001537//!!!! ALU SUBFAMILY SB2 WARNING ENT
RY !!!!!/0.0063:52:50//HOMO SAPIENS (HUMAN).//P391
91
F-HEM BB1001555//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/4.9e-23:69:63//HOMO SAPIENS (HUMAN).//P3918
8
F-HEM BB1001562//RABPHILIN-3A.//0.087:147:27//RATTU
S NORVEGICUS (RAT).//P47709
F-HEM BB1001564//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//5.9e-27:107:54//HOMO SAPIENS (HUMAN).//P08547
40 F-HEM BB1001565//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/1.9e-12:51:54//HOMO SAPIENS (HUMAN).//P391
94
F-HEM BB1001585
F-HEM BB1001586
F-HEM BB1001588//HYPOTHETICAL 12.3 KD PROTEIN IN GA
P1-NAP1 INTERGENIC REGION.//0.0031:31:48//SACCHARO
MYCES CEREVISIAE (BAKER'S YEAST).//P36140
F-HEM BB1001603
F-HEM BB1001618//RETROVIRUS-RELATED POL POLYPROTEIN
[CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); E

2925

NDONUCLEASE].//0.00076:47:44//MUS MUSCULUS (MOUSE).//P11369
 F-HEMBB1001619//SMALL PROLINE RICH PROTEIN II (SPR-II) (CLONE 930).//1.0:52:32//HOMO SAPIENS (HUMAN).//P22531
 F-HEMBB1001630
 F-HEMBB1001635//METALLOTHIONEIN-LIKE PROTEIN TYPE 2 A.//1.0:27:44//LYCOPERSICON ESCULENTUM (TOMATO).//Q40157
 F-HEMBB1001637//!!!! ALU SUBFAMILY J WARNING ENTRY 10
 !!!!!/0.0042:26:73//HOMO SAPIENS (HUMAN).//P39188
 F-HEMBB1001641
 F-HEMBB1001653//SURVIVAL MOTOR NEURON PROTEIN 1.//0.51:36:47//CANIS FAMILIARIS (DOG).//002771
 F-HEMBB1001665//HOMEBOX PROTEIN ENGRAILED-1 (HUN-1).//0.0030:135:34//HOMO SAPIENS (HUMAN).//Q05925
 F-HEMBB1001668//PROBABLE 60S RIBOSOMAL PROTEIN L39.//0.99:25:44//CAENORHABDITIS ELEGANS.//P52814
 F-HEMBB1001673//HYPOTHETICAL 46.1 KD PROTEIN IN ER 20
 P5-ORC6 INTERGENIC REGION.//0.0054:128:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38823
 F-HEMBB1001684//SUPPRESSOR PROTEIN SRP40.//0.56:81:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P32583
 F-HEMBB1001685//CYTOCHROME C OXIDASE POLYPEPTIDE V III-HEART PRECURSOR (EC 1.9.3.1) (VIIIB) (IX).//1.0:21:47//BOS TAURUS (BOVINE).//P10175
 F-HEMBB1001695//MYOSIN IC HEAVY CHAIN.//8.9e-05:86:40//ACANTHAMOEBA CASTELLANII (AMOEBA).//P10569
 F-HEMBB1001704//!!!! ALU SUBFAMILY SX WARNING ENTRY
 Y !!!!!/9.0e-08:35:71//HOMO SAPIENS (HUMAN).//P39195
 F-HEMBB1001706//CORNIFIN (SMALL PROLINE-RICH PROTEIN I) (SPR-I) (SMALL PROLINE-RICH SQUAMOUS CELL MARKER) (SPRP).//0.91:39:41//SUS SCROFA (PIC).//P3523
 F-HEMBB1001707//FERREDOXIN-LIKE PROTEIN IN NIF REGION.//1.0:43:23//BRADYRHIZOBIUM JAPONICUM.//P27394
 F-HEMBB1001717//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 4 (EC 1.6.5.3) (FRAGMENT).//1.0:71:25//LEMUR CATTIA (RING-TAILED LEMUR).//Q34878
 F-HEMBB1001735//!!!! ALU SUBFAMILY SQ WARNING ENTRY
 Y !!!!!/9.0e-35:97:74//HOMO SAPIENS (HUMAN).//P39194
 F-HEMBB1001736//EUKARYOTIC TRANSLATION INITIATION FACTOR 3 BETA SUBUNIT (EIF-3 BETA) (EIF3 P116) (EIF3 P110).//0.00069:180:28//HOMO SAPIENS (HUMAN).//P55884
 F-HEMBB1001747

2926

F-HEMBB1001749//!!!! ALU SUBFAMILY SX WARNING ENTRY
 Y !!!!!/1.8e-43:75:70//HOMO SAPIENS (HUMAN).//P39195
 F-HEMBB1001753//PROTEIN Q300.//0.00091:16:81//MUS MUSCULUS (MOUSE).//Q02722
 F-HEMBB1001756//CYCLIN-DEPENDENT KINASES REGULATOR Y SUBUNIT 2 (XE-P9).//0.94:35:42//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//Q91879
 F-HEMBB1001760
 F-HEMBB1001762//GENE 35 PROTEIN (GP35).//0.76:21:47//MYCOBACTERIOPHAGE L5.//Q05245
 F-HEMBB1001785
 F-HEMBB1001797//CHLOROPLAST 50S RIBOSOMAL PROTEIN L35.//0.99:41:31//PORPHYRA PURPUREA.//P51270
 F-HEMBB1001802
 F-HEMBB1001812//!!!! ALU SUBFAMILY SP WARNING ENTRY
 Y !!!!!/2.2e-39:54:77//HOMO SAPIENS (HUMAN).//P39193
 F-HEMBB1001816//!!!! ALU SUBFAMILY SQ WARNING ENTRY
 Y !!!!!/1.1e-19:97:57//HOMO SAPIENS (HUMAN).//P39194
 F-HEMBB1001831//HYPOTHETICAL 45.6 KD PROTEIN IN COX5A-ALC11 INTERGENIC REGION.//0.62:204:23//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53951
 F-HEMBB1001834//GLYCINE-RICH RNA-BINDING PROTEIN 1 (FRAGMENT).//0.0014:40:45//SORGHUM VULGARE (SORGHUM).//Q99069
 F-HEMBB1001836//!!!! ALU SUBFAMILY SB2 WARNING ENTRY
 Y !!!!!/7.1e-14:85:61//HOMO SAPIENS (HUMAN).//P39191
 F-HEMBB1001839//PROBABLE E4 PROTEIN.//0.61:49:34//HUMAN PAPILLOMAVIRUS TYPE 6C.//P20969
 F-HEMBB1001850
 F-HEMBB1001863//!!!! ALU SUBFAMILY SQ WARNING ENTRY
 Y !!!!!/2.7e-30:57:68//HOMO SAPIENS (HUMAN).//P39194
 F-HEMBB1001867
 F-HEMBB1001868//EXTENSIN PRECURSOR (CELL WALL HYDROXYPROLINE-RICH GLYCOPROTEIN).//0.00036:47:53//NICOTIANA TABACUM (COMMON TOBACCO).//P13983
 F-HEMBB1001869//!!!! ALU SUBFAMILY J WARNING ENTRY
 Y !!!!!/7.0e-11:95:45//HOMO SAPIENS (HUMAN).//P39188
 F-HEMBB1001872//HYPOTHETICAL 8.2 KD PROTEIN IN LEF8-FP INTERGENIC REGION.//1.0:34:38//AUTOGRAHA CALIFORNICA NUCLEAR POLYHEDROSIS VIRUS (ACMNPV).//P41459
 F-HEMBB1001874
 50 F-HEMBB1001875

- F-HEM BB1001880
 F-HEM BB1001899//GENE 11 PROTEIN. //1.0:45:31//SPIRO
 PLASMA VIRUS SPV1-R8A2B. //P15902
 F-HEM BB1001905//HYPOTHETICAL 81.7 KD PROTEIN IN MO
 L1-NAT2 INTERGENIC REGION. //8.8e-54:216:51//SACCHA
 ROMYCES CEREVISIAE (BAKER'S YEAST). //P48234
 F-HEM BB1001906
 F-HEM BB1001908//MONOCYTIC LEUKEMIA ZINC FINGER PRO
 TEIN. //6.3e-51:138:80//HOMO SAPIENS (HUMAN). //Q927
 94
 F-HEM BB1001910
 F-HEM BB1001911
 F-HEM BB1001915//UBIQUITIN CARBOXYL-TERMINAL HYDROL
 ASE 64E (EC 3.1.2.15) (UBIQUITIN THIOLESTERASE 64E)
 (UBIQUITIN-SPECIFIC PROCESSING PROTEASE 64E) (DEU
 BIQUITINATING ENZYME 64E). //2.3e-27:71:70//DROSOPH
 ILA MELANOGASTER (FRUIT FLY). //Q24574
 F-HEM BB1001921//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //9.8e-13:75:53//HOMO SAPIENS (HUMAN). //P08547
 F-HEM BB1001922
 F-HEM BB1001925//EPITHELIAL MEMBRANE PROTEIN-1 (EMP
 -1) (TUMOR-ASSOCIATED MEMBRANE PROTEIN). //1.0:55:30
 //MUS MUSCULUS (MOUSE). //P47801
 F-HEM BB1001930//HYPOTHETICAL 9.6 KD PROTEIN K10D2.
 7 IN CHROMOSOME III. //0.43:49:26//CAENORHABDITIS E
 LEGANS. //Q09412
 F-HEM BB1001944//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/5.1e-34:63:85//HOMO SAPIENS (HUMAN). //P391
 89
 F-HEM BB1001945//NONSPECIFIC LIPID-TRANSFER PROTEIN
 (LTP) (PHOSPHOLIPID TRANSFER PROTEIN) (PLTP). //0.
 28:45:40//AMARANTHUS CAUDATUS (LOVE-LIES-BLEEDING)
 (INCA-WHEAT). //P80450
 F-HEM BB1001947//PROTEIN UL24. //0.48:42:47//HERPES
 SIMPLEX VIRUS (TYPE 1/ STRAIN 17). //P10208
 F-HEM BB1001950//HYPOTHETICAL 42.6 KD PROTEIN IN GS
 HB-ANSB INTERGENIC REGION (0378). //1.6e-24:162:36/
 /ESCHERICHIA COLI. //P52062
 F-HEM BB1001952
 F-HEM BB1001953
 F-HEM BB1001957//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/2.7e-11:51:60//HOMO SAPIENS (HUMAN). //P3918
 8
 F-HEM BB1001962//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/7.6e-24:163:42//HOMO SAPIENS (HUMAN). //P391
 88
 F-HEM BB1001967//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/2.1e-35:55:80//HOMO SAPIENS (HUMAN). //P391
 89
 F-HEM BB1001973//!!!! ALU SUBFAMILY SC WARNING ENTR
- Y !!!!!/2.1e-37:108:75//HOMO SAPIENS (HUMAN). //P39
 192
 F-HEM BB1001983//LYSIS PROTEIN (E PROTEIN) (GPE). //
 0.84:45:37//BACTERIOPHAGE ALPHA-3. //P31280
 F-HEM BB1001988
 F-HEM BB1001990
 F-HEM BB1001996//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //1.4e-14:98:40//HOMO SAPIENS (HUMAN). //P08547
 F-HEM BB1001997//!!!! ALU SUBFAMILY J WARNING ENTRY
 10 !!!!!/4.1e-19:38:73//HOMO SAPIENS (HUMAN). //P3918
 8
 F-HEM BB1002002//CYTOCHROME C BIOGENESIS PROTEIN CC
 SA. //1.0:150:25//PORPHYRA PURPUREA. //P51369
 F-HEM BB1002005//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!/7.6e-12:94:40//HOMO SAPIENS (HUMAN). //P391
 95
 F-HEM BB1002009
 F-HEM BB1002015//HYPOTHETICAL 7.7 KD PROTEIN IN MRR
 -TSR INTERGENIC REGION(F67). //1.0:17:47//ESCHERICH
 20 IA COLI. //P39395
 F-HEM BB1002042//CYTOCHROME P450 4C1 (EC 1.14.14.1)
 (CYP1VC1). //2.4e-50:139:55//BLABERUS DISCOIDALIS
 (TROPICAL COCKROACH). //P29981
 F-HEM BB1002043//HYPOTHETICAL 9.5 KD PROTEIN IN DHF
 R 3' REGION (ORF3). //0.052:40:42//HERPESVIRUS SAIMI
 RI (SUBGROUP C / STRAIN 488). //P22577
 F-HEM BB1002044//CELLULOSE COMPLEMENTING PROTEIN. //
 0.45:87:33//ACETOBACTER XYLINUM (ACETOBACTER PASTE
 URIANUS). //P37697
 30 F-HEM BB1002045//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
 4) (AGL). //0.78:18:55//HOMO SAPIENS (HUMAN). //P039
 28
 F-HEM BB1002049
 F-HEM BB1002050//SALIVARY PROLINE-RICH PROTEIN PREC
 URSOR (CLONE CP7) [CONTAINS: BASIC PEPTIDE P-F] (F
 RAGMENT). //1.0e-06:188:27//HOMO SAPIENS (HUMAN). //
 P02812
 F-HEM BB1002068//HOMEODOMAIN PROTEIN HOX-A4 (CHOX-1.
 4). //0.0023:56:44//GALLUS GALLUS (CHICKEN). //P1727
 40 7
 F-HEM BB1002069//EXTENSIN PRECURSOR (CELL WALL HYDR
 OXYPROLINE-RICH GLYCOPROTEIN). //0.0074:134:33//NIC
 OTIANA TABACUM (COMMON TOBACCO). //P13983
 F-HEM BB1002092//ENV POLYPROTEIN PRECURSOR (COAT PO
 LYPROTEIN) [CONTAINS: OUTER MEMBRANE PROTEIN GP70;
 TRANSMEMBRANE PROTEIN P20E]. //2.4e-07:75:40//BABOO
 N ENDOGENOUS VIRUS (STRAIN M7). //P10269
 F-HEM BB1002094//!!!! ALU SUBFAMILY SB2 WARNING ENT
 RY !!!!!/1.9e-24:63:82//HOMO SAPIENS (HUMAN). //P39
 191

- F-HEMBB1002115//EC PROTEIN HOMOLOG (ZINC-METALLOTHIONEIN CLASS II).//0.94:26:42//ZEA MAYS (MAIZE).//P43401
- F-HEMBB1002134//ZINC-FINGER PROTEIN NEURO-D4.//4.6e-57:176:67//RATTUS NORVEGICUS (RAT).//P56163
- F-HEMBB1002139//CHLOROPLAST 50S RIBOSOMAL PROTEIN L35.//1.0:17:52//PORPHYRA PURPUREA.//P51270
- F-HEMBB1002142//EARLY NODULIN 20 PRECURSOR (N-20).//0.087:52:36//MEDICAGO TRUNCATULA (BARREL MEDIC).//P93329
- F-HEMBB1002152//HYPOTHETICAL 12.3 KD PROTEIN IN RP L3-RPL33 INTERGENIC REGION (ORF102).//5.8e-05:61:37//CYANOPHORA PARADOXA.//P15811
- F-HEMBB1002189//HYPOTHETICAL PROTEIN UL125.//1.0:77:32//HUMAN CYTOMEGALOVIRUS (STRAIN AD169).//P16835
- F-HEMBB1002190
- F-HEMBB1002193//TYROSINE-PROTEIN KINASE RECEPTOR TYRO3 PRECURSOR (TYROSINE-PROTEIN KINASE RSE) (TYROSINE-PROTEIN KINASE SKY) (TYROSINE-PROTEIN KINASE DTK).//1.2e-27:59:100//HOMO SAPIENS (HUMAN).//Q06418
- F-HEMBB1002217//ZINC FINGER PROTEIN 184 (FRAGMENT).//6.6e-22:106:50//HOMO SAPIENS (HUMAN).//Q99676
- F-HEMBB1002218//PROTEIN Q300.//0.85:19:52//MUS MUSCULUS (MOUSE).//Q02722
- F-HEMBB1002232//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//9.6e-21:56:71//HOMO SAPIENS (HUMAN).//P39195
- F-HEMBB1002247
- F-HEMBB1002249//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//7.2e-29:93:69//HOMO SAPIENS (HUMAN).//P39194
- F-HEMBB1002254//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//2.0e-29:101:67//HOMO SAPIENS (HUMAN).//P39194
- F-HEMBB1002255//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 3 (EC 1.6.5.3).//1.0:73:28//PARAMECIUM TETRAURELIA.//P15579
- F-HEMBB1002266//GLUTAMIC ACID-RICH PROTEIN PRECURSOR.//0.0079:151:26//PLASMODIUM FALCIPARUM (ISOLATE FC27 / PAPUA NEW GUINEA).//P13816
- F-HEMBB1002280//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//5.2e-15:182:36//NYCTICEBUS COUCANG (SLOW LORIS).//P08548
- F-HEMBB1002300
- F-HEMBB1002306//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//0.00011:26:84//HOMO SAPIENS (HUMAN).//P39195
- F-HEMBB1002327//!!!! ALU SUBFAMILY SB WARNING ENTRY
- Y !!!!!//4.1e-11:41:85//HOMO SAPIENS (HUMAN).//P39189
- F-HEMBB1002329//HYPOTHETICAL 74.0 KD PROTEIN IN CAJ1-HOM3 INTERGENIC REGION.//9.9e-17:232:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40032
- F-HEMBB1002340
- F-HEMBB1002342//HYPOTHETICAL 32.5 KD PROTEIN IN MSB6-BMH2 INTERGENIC REGION.//3.6e-40:102:57//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q03835
- 10 F-HEMBB1002358//THYMIDYLATE KINASE (EC 2.7.4.9) (DTMP KINASE).//6.1e-30:63:96//HOMO SAPIENS (HUMAN).//P23919
- F-HEMBB1002359//HYPOTHETICAL 7.1 KD PROTEIN C6G9.01C IN CHROMOSOME I.//0.97:28:46//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q92346
- F-HEMBB1002364//RETROVIRUS-RELATED POLYPROTEIN (FRAGMENT).//0.47:119:25//HOMO SAPIENS (HUMAN).//P12895
- F-HEMBB1002371//HYPOTHETICAL 15.5 KD PROTEIN C2F7.12 IN CHROMOSOME I PRECURSOR.//3.0e-05:111:30//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09703
- F-HEMBB1002381//PUTATIVE CUTICLE COLLAGEN C09G5.4.//0.34:105:34//CAENORHABDITIS ELEGANS.//Q09455
- F-HEMBB1002383//ATP SYNTHASE A CHAIN (EC 3.6.1.34) (PROTEIN 6).//0.049:103:32//AQUIFEX AEOLICUS.//Q066566
- F-HEMBB1002387//10 KD CHAPERONIN (PROTEIN CPN10) (PROTEIN GROES) (HEAT SHOCK PROTEIN 11).//0.18:75:28//RICKETTSIA TSUTSUGAMUSHI.//P16626
- 30 F-HEMBB1002409//HIGH MOBILITY GROUP PROTEIN HMG-Y.//0.014:61:36//MUS MUSCULUS (MOUSE).//P17095
- F-HEMBB1002415
- F-HEMBB1002425//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//3.8e-18:55:70//HOMO SAPIENS (HUMAN).//P39194
- F-HEMBB1002442//LIN-10 PROTEIN.//5.1e-15:121:31//CAENORHABDITIS ELEGANS.//P34692
- F-HEMBB1002453//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//1.2e-32:54:75//HOMO SAPIENS (HUMAN).//P39189
- F-HEMBB1002457//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.5e-07:31:64//HOMO SAPIENS (HUMAN).//P39188
- F-HEMBB1002458//MALE SPECIFIC SPERM PROTEIN MST84DA.//0.92:28:53//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01642
- F-HEMBB1002477//DNA-DIRECTED RNA POLYMERASE II LARGE SUBUNIT (EC 2.7.7.6) (RPB1) (FRAGMENT).//0.0066:198:27//CRICETULUS GRISEUS (CHINESE HAMSTER).//P11414

2931

F-HEMBB1002489//SPLICEOSOME ASSOCIATED PROTEIN 49 (SAP 49) (SF3B53).//0.030:182:28//HOMO SAPIENS (HUMAN).//Q15427

F-HEMBB1002492

F-HEMBB1002495//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!//2.1e-08:41:75//HOMO SAPIENS (HUMAN).//P39192

F-HEMBB1002502//RETROVIRUS-RELATED POL POLYPROTEIN (FRAGMENT).//0.00030:31:77//HOMO SAPIENS (HUMAN).//P12895

F-HEMBB1002509

F-HEMBB1002510

F-HEMBB1002520//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//6.8e-36:162:50//NYCTICEBUS COUCANG (SLOW LORIS).//P08548

F-HEMBB1002522//7 KD PROTEIN (ORF 4).//0.77:32:40//CHRYSANTHEMUM VIRUS B(CVB).//P37990

F-HEMBB1002531

F-HEMBB1002534//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//3.1e-36:80:73//HOMO SAPIENS (HUMAN).//P39195

F-HEMBB1002545

F-HEMBB1002550//HOMEODOMAIN PROTEIN HOX-D11 (HOX-4.6) (HOX-5.5).//3.8e-05:83:34//MUS MUSCULUS (MOUSE).//P23813

F-HEMBB1002556

F-HEMBB1002579//SPLICING FACTOR U2AF 35 KD SUBUNIT (U2 AUXILIARY FACTOR 35 KD SUBUNIT) (U2 SNRNP AUXILIARY FACTOR SMALL SUBUNIT) (FRAGMENT).//5.0e-06:27:77//SUS SCROFA (PIG).//Q29350

F-HEMBB1002582//PROTEINASE INHIBITOR.//1.0:27:40//SOLANUM MELONGENA (EGGPLANT) (AUBERGINE).//P01078

F-HEMBB1002590//HYPOTHETICAL PROTEIN IN MMSB 3' REGION (ORF1) (FRAGMENT).//1.9e-20:90:54//PSEUDOMONAS AERUGINOSA.//P28812

F-HEMBB1002596

F-HEMBB1002600//NOVEL ANTIGEN 2 (NAG-2).//1.9e-60:187:59//HOMO SAPIENS (HUMAN).//O14817

F-HEMBB1002601//M PROTEIN, SEROTYPE 6 PRECURSOR.//1.0:71:35//STREPTOCOCCUS PYOGENES.//P08089

F-HEMBB1002603

F-HEMBB1002607//SALIVARY PROLINE-RICH PROTEIN P0 (ALLELE K) [CONTAINS: PEPTIDE P-D] (FRAGMENT).//0.0032:142:33//HOMO SAPIENS (HUMAN).//P10162

F-HEMBB1002610//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//2.0e-11:79:49//HOMO SAPIENS (HUMAN).//P08547

F-HEMBB1002613//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//3.9e-08:41:60//HOMO SAPIENS (HUMAN).//P39188

F-HEMBB1002614//HYPOTHETICAL 9.5 KD PROTEIN.//1.0: 50

2932

40:35//VACCINIA VIRUS(STRAIN COPENHAGEN).//P20553

F-HEMBB1002617//INSECT TOXIN 1 (BOT IT1).//1.0:44:29//BUTHUS OCCITANUS TUNETANUS (COMMON EUROPEAN SCORPION).//P55902

F-HEMBB1002623//HYPOTHETICAL 9.7 KD PROTEIN (ORF88) (PUTATIVE DNA-BINDING PROTEIN).//0.42:31:54//BACTERIOPHAGE P4.//P12552

F-HEMBB1002635//STRESS-ACTIVATED PROTEIN KINASE JNK3 (EC 2.7.1.-) (C-JUNN-TERMINAL KINASE 3) (MAP KINASE P49 3F12).//6.2e-17:44:95//HOMO SAPIENS (HUMAN).//P53779

F-HEMBB1002664//SMALL NUCLEAR RIBONUCLEOPROTEIN ASSOCIATED PROTEIN B (SM-B) (SNRNP-B) (SM11) (FRAGMENT).//1.0:57:36//RATTUS NORVEGICUS (RAT).//P17136

F-HEMBB1002677//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//1.9e-06:194:34//NYCTICEBUS COUCANG (SLOW LORIS).//P08548

F-HEMBB1002683//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 5 (EC 1.6.5.3) (FRAGMENT).//0.96:56:35//LEMUR CATTAL (RING-TAILED LEMUR).//Q34879

F-HEMBB1002684//SILLUCIN.//1.0:16:50//RHIZOMUCOR PUSILLUS.//P02885

F-HEMBB1002686

F-HEMBB1002692

F-HEMBB1002697//HELIX-DESTABILIZING PROTEIN (SINGLE-STRANDED DNA BINDING PROTEIN) (GPV).//0.57:36:38//BACTERIOPHAGE FD, BACTERIOPHAGE F1, AND BACTERIOPHAGE M13.//P03669

F-HEMBB1002699

30 F-HEMBB1002702

F-HEMBB1002705//HYPOTHETICAL 34.8 KD PROTEIN C4H3.04C IN CHROMOSOME I.//3.6e-40:180:37//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q10212

F-HEMBB1002712

F-MAMMA1000009//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//1.2e-32:95:75//HOMO SAPIENS (HUMAN).//P39189

F-MAMMA1000019

40 F-MAMMA1000020//DIMETHYLANILINE MONOOXYGENASE [N-OXIDE FORMING] 5 (EC 1.14.13.8) (HEPATIC FLAVIN-CONTAINING MONOOXYGENASE 5) (FMO 5) (DIMETHYLANILINE OXIDASE 5).//5.2e-12:24:100//HOMO SAPIENS (HUMAN).//P49326

F-MAMMA1000025//BETA-2-MICROGLOBULIN PRECURSOR.//1.0:73:26//BRACHYDANIORERI (ZEBRAFISH) (ZEBRA DANIO).//Q04475

F-MAMMA1000043//HYPOTHETICAL PXBL-1 PROTEIN (FRAGMENT).//0.057:130:31//BOVINE LEUKEMIA VIRUS (JAPANESE ISOLATE BLV-1) (BLV).//P03412

F-MAMMA1000045

2933

F-MAMMA1000055//TESTIN 2 (TES2) [CONTAINS: TESTIN 1 (TES1)].//7.5e-44:138:55//MUS MUSCULUS (MOUSE).//P47226

F-MAMMA1000057//!!!! ALU SUBFAMILY SQ WARNING ENTR Y !!!!!/1.2e-39:92:69//HOMO SAPIENS (HUMAN).//P39194

F-MAMMA1000069//HYPOTHETICAL 29.3 KD PROTEIN (ORF92).//0.0044:96:34//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPMNPV).//010341

F-MAMMA1000084//!!!! ALU SUBFAMILY SX WARNING ENTR Y !!!!!/5.4e-28:94:73//HOMO SAPIENS (HUMAN).//P39195

F-MAMMA1000085//PUTATIVE CYSTEINYL-TRNA SYNTHETASE C29E6.06C (EC 6.1.1.16) (CYSTEINE-- TRNA LIGASE) (CYSRS).//6.6e-38:90:51//SCHIZOSACCHAROMYCESPOMBE (FISSION YEAST).//Q09860

F-MAMMA1000092//!!!! ALU SUBFAMILY SC WARNING ENTR Y !!!!!/6.4e-30:43:86//HOMO SAPIENS (HUMAN).//P39192

F-MAMMA1000103//SMALL PROLINE RICH PROTEIN II (SPR-II) (CLONE 930).//0.038:17:52//HOMO SAPIENS (HUMAN).//P22531

F-MAMMA1000117//50S RIBOSOMAL PROTEIN L24E (HL21/H L22).//0.90:25:48//HALOARcula MARISMORTUI (HALOBACTERIUM MARISMORTUI).//P14116

F-MAMMA1000129//HYPOTHETICAL BHLF1 PROTEIN.//0.0016:75:40//EPSTEIN-BARRVIRUS (STRAIN B95-8) (HUMAN HERPESVIRUS 4).//P03181

F-MAMMA1000133

F-MAMMA1000134//HYPOTHETICAL PROTEIN MJ0647.//1.0:41:41//METHANOCOCCUS JANNASCHII.//Q58063

F-MAMMA1000139//GUANINE NUCLEOTIDE-BINDING PROTEIN G(I)/G(S)/G(O) GAMMA-3 SUBUNIT.//0.99:69:28//BOSTAURUS (BOVINE), AND MUS MUSCULUS (MOUSE).//P29798

F-MAMMA1000143//CALPAIN INHIBITOR (CALPASTATIN) (FRAGMENT).//0.023:111:27//MUS MUSCULUS (MOUSE).//P51125

F-MAMMA1000155//PUTATIVE CUTICLE COLLAGEN C09G5.5.//0.018:125:34//CAENORHABDITIS ELEGANS.//Q09456

F-MAMMA1000163//MERCURIC TRANSPORT PROTEIN PERIPLASMIC COMPONENT PRECURSOR (PERIPLASMIC MERCURY ION BINDING PROTEIN) (MERCURY SCAVENGER PROTEIN).//0.11:88:25//SHEWANELLA PUTREFACIENS (PSEUDOMONAS PUTREFACIENS).//Q54463

F-MAMMA1000171

F-MAMMA1000173//DREBRIN E.//7.6e-41:197:43//HOMO SAPIENS (HUMAN).//Q16643

F-MAMMA1000175//GAMMA-THIONIN HOMOLOG PPT PRECURSOR.//0.92:39:38//PETUNIA INTEGRIFOLIA (VIOLET-FLOWERED PETUNIA) (PETUNIA INFLATA).//Q40901

2934

F-MAMMA1000183//ZINC FINGER PROTEIN 84 (ZINC FINGER PROTEIN HPF2).//2.4e-106:249:61//HOMO SAPIENS (HUMAN).//P51523

F-MAMMA1000198//MALE SPECIFIC SPERM PROTEIN MST84D.//0.0014:35:42//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01645

F-MAMMA1000221

F-MAMMA1000227//6.8 KD MITOCHONDRIAL PROTEOLIPID.//1.0:30:40//MUS MUSCULUS (MOUSE).//P56379

10 F-MAMMA1000241//PHOTOSYSTEM I REACTION CENTRE SUBUNIT X (PSI-K).//1.0:40:37//PORPHYRA PURPUREA.//P51370

F-MAMMA1000251//HYPOTHETICAL 6.8 KD PROTEIN IN FIC-PPIA INTERGENIC REGION.//0.99:29:48//SALMONELLA TYPHIMURIUM.//P37771

F-MAMMA1000254//HYPOTHETICAL 6.0 KD PROTEIN IN THI 12 5' REGION.//1.0:20:50//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53820

F-MAMMA1000257//HYPOTHETICAL 50.0 KD PROTEIN IN HEML 3' REGION (ORF2).//0.22:50:44//PSEUDOMONAS AERUGINOSA.//Q51470

F-MAMMA1000264//GASTRIN-RELEASING PEPTIDE RECEPTOR (GRP-R) (GRP-PREFERRING BOMBESIN RECEPTOR).//0.80:39:43//HOMO SAPIENS (HUMAN).//P30550

F-MAMMA1000266

F-MAMMA1000270//!!!! ALU SUBFAMILY SB WARNING ENTR Y !!!!!/9.5e-42:95:84//HOMO SAPIENS (HUMAN).//P39189

F-MAMMA1000277//PROCOLLAGEN ALPHA 1(II) CHAIN PRECURSOR [CONTAINS: CHONDROCALCIN].//0.0062:90:34//MUS MUSCULUS (MOUSE).//P28481

F-MAMMA1000278//C-HORDEIN (CLONE PC HOR1-3) (FRAGMENT).//0.00096:59:33//HORDEUM VULGARE (BARLEY).//P17991

F-MAMMA1000279//!!!! ALU SUBFAMILY SX WARNING ENTR Y !!!!!/8.4e-17:56:76//HOMO SAPIENS (HUMAN).//P39195

F-MAMMA1000284//ARYL HYDROCARBON RECEPTOR NUCLEAR TRANSLOCATOR 2 (ARNT PROTEIN 2).//0.017:146:30//MUS MUSCULUS (MOUSE).//Q61324

F-MAMMA1000287//!!!! ALU SUBFAMILY SB WARNING ENTR Y !!!!!/1.5e-32:84:58//HOMO SAPIENS (HUMAN).//P39189

F-MAMMA1000302//C-HORDEIN (CLONE PC-919) (FRAGMENT).//1.0:42:33//HORDEUM VULGARE (BARLEY).//P17992

F-MAMMA1000307//PROBABLE E4 PROTEIN.//0.21:71:30//RHESUS PAPILLOMAVIRUS TYPE 1 (RHPV 1).//P24832

F-MAMMA1000309//COLLAGEN ALPHA 1(VIII) CHAIN PRECURSOR (ENDOTHELIAL COLLAGEN).//0.0026:141:36//HOMO SAPIENS (HUMAN).//P27658

F-MAMMA1000312

F-MAMMA1000313//DNA REPAIR PROTEIN RAD51 HOMOLOG (25 KDa PROTEIN) (FRAGMENT).//0.76:52:32//STAPHYLOCOCCUS AUREUS.//P31337

F-MAMMA1000331

F-MAMMA1000339//50S RIBOSOMAL PROTEIN L29P.//0.78:32:46//METHANOBACTERIUM THERMAUTOTROPHICUM.//026117

F-MAMMA1000340//HYPOTHETICAL 29.4 KDa PROTEIN IN STAG-LOS1 INTERGENIC REGION.//1.0:29:58//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P36039

F-MAMMA1000348//!!!! ALU SUBFAMILY J WARNING ENTRY!!!!//7.5e-09:63:60//HOMO SAPIENS (HUMAN).//P39188

F-MAMMA1000356//!!!! ALU SUBFAMILY J WARNING ENTRY!!!!//3.3e-05:42:52//HOMO SAPIENS (HUMAN).//P39188

F-MAMMA1000360

F-MAMMA1000361//!!!! ALU SUBFAMILY SB WARNING ENTRY!!!!//4.4e-33:84:72//HOMO SAPIENS (HUMAN).//P39189

F-MAMMA1000372//!!!! ALU SUBFAMILY SP WARNING ENTRY!!!!//6.6e-21:53:71//HOMO SAPIENS (HUMAN).//P39193

F-MAMMA1000385

F-MAMMA1000388//OX40L RECEPTOR PRECURSOR (ACT35 ANTIGEN) (TAX-TRANSCRIPTIONALLY ACTIVATED GLYCOPROTEIN 1 RECEPTOR) (CD134 ANTIGEN).//0.40:72:36//HOMO SAPIENS (HUMAN).//P43489

F-MAMMA1000395//RABPHILIN-3A (FRAGMENT).//0.032:125:25//MUS MUSCULUS (MOUSE).//P47708

F-MAMMA1000402//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//3.1e-28:266:40//HOMO SAPIENS (HUMAN).//P08547

F-MAMMA1000410//NADH-UBIQUINONE OXIDOREDUCTASE 13 KDa-B SUBUNIT (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-13KDa-B) (CI-13KDa-B) (B13).//5.9e-06:32:68//HOMO SAPIENS (HUMAN).//Q16718

F-MAMMA1000413//RETROVIRUS-RELATED POLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); ENDONUCLEASE].//6.7e-05:93:31//MUS MUSCULUS (MOUSE).//P11369

F-MAMMA1000414

F-MAMMA1000416//HYPOTHETICAL 32.0 KDa PROTEIN C09F5.2 IN CHROMOSOME III.//4.1e-28:119:53//CAENORHABDITIS ELEGANS.//Q09232

F-MAMMA1000421//!!!! ALU SUBFAMILY SQ WARNING ENTRY!!!!//2.7e-23:68:76//HOMO SAPIENS (HUMAN).//P39194

F-MAMMA1000422//METALLOTHIONEIN (MT).//0.037:42:42//GADUS MORHUA (ATLANTIC COD).//P51902

F-MAMMA1000423

F-MAMMA1000424//!!!! ALU SUBFAMILY SB WARNING ENTRY!!!!//0.048:23:73//HOMO SAPIENS (HUMAN).//P39189

F-MAMMA1000429//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS5.//2.7e-05:110:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q92331

F-MAMMA1000431//!!!! ALU SUBFAMILY SQ WARNING ENTRY!!!!//9.4e-15:85:58//HOMO SAPIENS (HUMAN).//P39194

F-MAMMA1000444//!!!! ALU SUBFAMILY SQ WARNING ENTRY!!!!//4.3e-25:65:76//HOMO SAPIENS (HUMAN).//P39194

F-MAMMA1000446//ZYXIN.//0.79:155:29//CALLUS CALLUS (CHICKEN).//Q04584

F-MAMMA1000458//HYPOTHETICAL 37.7 KDa PROTEIN C18B1.06 IN CHROMOSOME I.//0.0048:46:43//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09713

F-MAMMA1000468//PERIOD CLOCK PROTEIN (FRAGMENT).//0.50:20:55//DROSOPHILAROBUSTA (FRUIT FLY).//Q03296

F-MAMMA1000472//!!!! ALU SUBFAMILY J WARNING ENTRY!!!!//1.5e-17:106:55//HOMO SAPIENS (HUMAN).//P39188

F-MAMMA1000478//!!!! ALU SUBFAMILY SX WARNING ENTRY!!!!//3.9e-35:80:68//HOMO SAPIENS (HUMAN).//P39195

F-MAMMA1000483//!!!! ALU SUBFAMILY SP WARNING ENTRY!!!!//2.8e-24:74:77//HOMO SAPIENS (HUMAN).//P39193

F-MAMMA1000490//TYROSINE-PROTEIN KINASE TXK (EC 2.7.1.112) (PTK-RL-18) (RESTING LYMPHOCYTE KINASE).//0.43:21:57//MUS MUSCULUS (MOUSE).//P42682

F-MAMMA1000500//TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN).//0.61:33:54//HUMAN IMMUNODEFICIENCY VIRUS TYPE 1 (Z2/CDC-234 ISOLATE) (HIV-1).//P12506

F-MAMMA1000501//!!!! ALU SUBFAMILY SQ WARNING ENTRY!!!!//1.1e-32:43:83//HOMO SAPIENS (HUMAN).//P39194

F-MAMMA1000516

F-MAMMA1000522//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//0.0015:113:32//HOMO SAPIENS (HUMAN).//P08547
F-MAMMA1000524//HYPOTHETICAL HOST RANGE 8.5 KDa PROTEIN.//1.0:63:31//VACCINIA VIRUS (STRAIN WR).//P17359

F-MAMMA1000559//METALLOTHIONEIN-I (MT-I) (MT-IB/MT-IA).//0.31:16:50//CALLINECTES SAPIDUS (BLUE CRAB).//P55949

F-MAMMA1000565//FERREDOXIN-TYPE PROTEIN NAPP.//0.98:37:35//ESCHERICHIA COLI.//P33939

50 F-MAMMA1000567//!!!! ALU SUBFAMILY SX WARNING ENTRY

2937

Y !!!!!/5.5e-37:95:76//HOMO SAPIENS (HUMAN).//P391
95
F-MAMMA1000576//!!!! ALU SUBFAMILY SB2 WARNING ENT
RY !!!!!/4.1e-07:34:64//HOMO SAPIENS (HUMAN).//P39
191
F-MAMMA1000583
F-MAMMA1000585//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/1.5e-28:89:75//HOMO SAPIENS (HUMAN).//P391
94
F-MAMMA1000594//!!!! ALU SUBFAMILY SX WARNING ENTR 10
Y !!!!!/4.8e-24:38:71//HOMO SAPIENS (HUMAN).//P391
95
F-MAMMA1000597//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/1.1e-25:74:77//HOMO SAPIENS (HUMAN).//P391
95
F-MAMMA1000605//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/2.1e-18:83:50//HOMO SAPIENS (HUMAN).//P391
95
F-MAMMA1000612//HYPOTHETICAL 34.0 KD TRP-ASP REPEA
TS CONTAINING PROTEININ SIS1-MRPL2 INTERGENIC REGI 20
ON.//4.0e-42:166:48//SACCHAROMYCES CEREVISIAE (BAK
ER'S YEAST).//P41318
F-MAMMA1000616
F-MAMMA1000621
F-MAMMA1000623//METALLOTHIONEIN-1K (MT-1K).//0.004
5:25:48//HOMO SAPIENS(HUMAN).//P80296
F-MAMMA1000625//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T).//0.00078:79:35//MUS MUSCULUS (MOUSE).//P05143
F-MAMMA1000643//HYPOTHETICAL 9.3 KD PROTEIN.//1.0:
25:28//MAGUARI VIRUS.//P16607
F-MAMMA1000664
F-MAMMA1000669//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//4.2e-05:186:30//HOMO SAPIENS (HUMAN).//P08547
F-MAMMA1000670//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T).//1.6e-06:195:30//MUS MUSCULUS (MOUSE).//P05143
F-MAMMA1000672//VITELLOGENIC CARBOXYPEPTIDASE PREC
URSOR (EC 3.4.16.-).//3.8e-28:184:35//AEDES AEGYPT
I (YELLOWFEVER MOSQUITO).//P42660
F-MAMMA1000684//DNA-BINDING PROTEIN (VMW21).//1.1e
-07:55:56//HERPES SIMPLEX VIRUS (TYPE 1 / STRAIN 1 40
7).//P04487
F-MAMMA1000696//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/1.2e-31:97:74//HOMO SAPIENS (HUMAN).//P391
94
F-MAMMA1000707//METALLOTHIONEIN-II (MT-II) (MT-IIB
/MT-IIA).//0.31:19:42//CALLINECTES SAPIDUS (BLUE C
RAB).//P55950
F-MAMMA1000713//XYLOULOSE KINASE (EC 2.7.1.17) (XYL
ULOXINASE).//1.6e-05:88:35//LACTOBACILLUS PENTOSU
S.//P21939

2938

F-MAMMA1000714//PROTEIN-LYSINE 6-OXIDASE PRECURSOR
(EC 1.4.3.13) (LYSYLOXIDASE).//0.44:126:30//RATTU
S NORVEGICUS (RAT).//P16636
F-MAMMA1000718//METALLOTHIONEIN-II (MT-2E).//1.0:
51:31//ORYCTOLAGUS CUNICULUS (RABBIT).//P80292
F-MAMMA1000720//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/3.3e-28:60:71//HOMO SAPIENS (HUMAN).//P391
93
F-MAMMA1000723//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//3.7e-14:63:53//HOMO SAPIENS (HUMAN).//P08547
F-MAMMA1000731//CHROMODOMAIN-HELICASE-DNA-BINDING
PROTEIN 2 (CHD-2).//1.8e-43:258:43//HOMO SAPIENS
(HUMAN).//014647
F-MAMMA1000732//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/9.9e-12:76:55//HOMO SAPIENS (HUMAN).//P3918
8
F-MAMMA1000733
F-MAMMA1000734//NPL1 PROTEIN (SEC63 PROTEIN).//2.5
e-18:181:39//SACCHAROMYCES CEREVISIAE (BAKER'S YEA
ST).//P14906
F-MAMMA1000738//HYPOTHETICAL 116.5 KD PROTEIN C20G
8.09C IN CHROMOSOME I.//5.4e-52:196:58//SCHIZOSACC
HAROMYCES POMBE (FISSION YEAST).//P87115
F-MAMMA1000744//!!!! ALU SUBFAMILY SB1 WARNING ENT
RY !!!!!/6.3e-36:144:47//HOMO SAPIENS (HUMAN).//P3
9190
F-MAMMA1000746
F-MAMMA1000752
F-MAMMA1000760//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/6.6e-29:75:72//HOMO SAPIENS (HUMAN).//P391
95
F-MAMMA1000761//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/3.6e-09:59:64//HOMO SAPIENS (HUMAN).//P391
94
F-MAMMA1000775
F-MAMMA1000776//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/3.3e-35:99:74//HOMO SAPIENS (HUMAN).//P391
93
F-MAMMA1000778//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/2.1e-19:65:70//HOMO SAPIENS (HUMAN).//P391
95
F-MAMMA1000782
F-MAMMA1000798//HYPOTHETICAL PROTEIN ORF-1137.//0.
015:59:37//MUS MUSCULUS (MOUSE).//P11260
F-MAMMA1000802//MYOSIN IC HEAVY CHAIN.//0.35:94:41
//ACANTHAMOEBA CASTELLANII (AMOEBA).//P10569
F-MAMMA1000824//ACTIN 1.//0.046:60:31//ZEA MAYS (M
AIZE).//P02582
F-MAMMA1000831//PROBABLE NI/FE-HYDROGENASE 1 B-TYP
E CYTOCHROME SUBUNIT.//1.0:30:46//ESCHERICHIA COL

I.//P19929
 F-MAMMA1000839//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/1.1e-28:80:58//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1000841//PUTATIVE AMIDASE (EC 3.5.1.4).//1.
 5e-39:130:36//METHANOBACTERIUM THERMOAUTOTROPHICU
 M.//027540
 F-MAMMA1000842//C-HORDEIN (CLONE PC-919) (FRAGMEN
 T).//0.064:43:41//HORDEUM VULGARE (BARLEY).//P1799
 2
 F-MAMMA1000843
 F-MAMMA1000845//NADH-UBIQUINONE OXIDOREDUCTASE CHA
 IN 2 (EC 1.6.5.3).//0.43:58:34//DROSOPHILA YAKUBA
 (FRUIT FLY).//P03895
 F-MAMMA1000851//CUTICLE COLLAGEN 34.//0.019:107:29
 //CAENORHABDITIS ELEGANS.//P34687
 F-MAMMA1000855//SPLICEOSOME ASSOCIATED PROTEIN 62
 (SAP 62) (SF3A66).//0.00098:149:32//HOMO SAPIENS
 (HUMAN).//Q15428
 F-MAMMA1000856//METALLOTHIONEIN (MT).//0.63:39:41/
 /POTAMON POTAMIOS.//P55952
 F-MAMMA1000859//GLYCOPROTEIN X PRECURSOR.//0.014:1
 92:28//EQUINE HERPESVIRUS TYPE 1 (STRAIN AB4P) (EH
 V-1).//P28968
 F-MAMMA1000862//DISINTEGRIN KISTRIN (PLATELET AGGR
 EGATION ACTIVATION INHIBITOR).//1.0:66:27//AGKISTR
 ODN RHODOSTOMA (MALAYAN PIT VIPER) (CALLOSELASMA
 RHODOSTOMA).//P17494
 F-MAMMA1000863//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/5.4e-16:41:68//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1000865//SALIVARY PROLINE-RICH PROTEIN II-1
 (FRAGMENT).//0.030:100:32//HOMO SAPIENS (HUMAN)./
 /P81489
 F-MAMMA1000867//APTOTOXIN IX (PARALYTIC PEPTIDE I
 X) (PP IX).//0.98:43:32//APTOSTICHUS SCHLINGERI (T
 RAP-DOOR SPIDER).//P49272
 F-MAMMA1000875//PROLINE-RICH PEPTIDE P-B.//0.18:2
 1:47//HOMO SAPIENS (HUMAN).//P02814
 F-MAMMA1000876//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/1.5e-22:85:71//HOMO SAPIENS (HUMAN).//P391
 89
 F-MAMMA1000877//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/1.2e-38:62:74//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1000880//COLLAGEN ALPHA 1(XII) CHAIN (FRAGM
 ENTS).//0.49:79:32//BOS TAURUS (BOVINE).//P25508
 F-MAMMA1000883//HYPOTHETICAL 6.1 KD PROTEIN C03B1.
 10 IN CHROMOSOME X.//0.87:15:60//CAENORHABDITIS EL
 EGANS.//Q11116

F-MAMMA1000897//INTER-ALPHA-TRYPSIN INHIBITOR HEAV
 Y CHAIN H4 PRECURSOR (ITI HEAVY CHAIN H4) (INTER-A
 LPHA-TRYPSIN INHIBITOR FAMILY HEAVY CHAIN-RELATED
 PROTEIN) (PLASMA KALLIKREIN SENSITIVE GLYCOPROTEIN
 120) (PK-120).//5.3e-17:130:40//HOMO SAPIENS (HUM
 AN).//Q14624
 F-MAMMA1000905
 F-MAMMA1000906
 F-MAMMA1000908//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG.//8.0e-17:70:62//HOMO SAPIENS (HUMAN).//P08547
 10
 F-MAMMA1000914//HYPOTHETICAL 6.2 KD PROTEIN.//0.9
 7:36:36//THERMOPROTEUSTENAX VIRUS 1 (STRAIN KRA1)
 (TTV1).//P19299
 F-MAMMA1000921
 F-MAMMA1000931//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/5.6e-10:49:65//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1000940//MITOCHONDRIAL 60S RIBOSOMAL PROTEI
 N L32.//0.42:22:54//RECLINOMONAS AMERICANA.//02128
 1
 F-MAMMA1000941//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/2.3e-25:55:69//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1000942//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!/3.3e-08:36:75//HOMO SAPIENS (HUMAN).//P391
 94
 F-MAMMA1000943
 F-MAMMA1000956//SMALL HISTIDINE-ALANINE-RICH PROTE
 IN PRECURSOR (SHARP) (ANTIGEN 57).//0.041:122:25//
 PLASMODIUM FALCIPARUM (ISOLATE FC27 / PAPUAN NEW GUI
 NEA).//P04930
 F-MAMMA1000957//HEAT-STABLE ENTEROTOXIN A2 PRECURS
 OR (STA2).//0.024:37:37//ESCHERICHIA COLI.//Q47185
 F-MAMMA1000962//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/6.0e-39:61:78//HOMO SAPIENS (HUMAN).//P391
 89
 F-MAMMA1000968//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!/0.0054:29:72//HOMO SAPIENS (HUMAN).//P3919
 4
 F-MAMMA1000975//CUTICLE COLLAGEN DPY-2 PRECURSOR./
 /1.0:93:30//CAENORHABDITIS ELEGANS.//P35799
 F-MAMMA1000979//PROLINE-RICH PEPTIDE P-B.//0.012:1
 2:66//HOMO SAPIENS (HUMAN).//P02814
 F-MAMMA1000987//HYPOTHETICAL PROTEIN LAMBDA-SP34./
 /1.0:47:40//MUS MUSCULUS (MOUSE).//P15973
 F-MAMMA1000998
 F-MAMMA1001003//PROBABLE E5 PROTEIN.//1.0:52:42//H
 UMAN PAPILOMAVIRUS TYPE 33.//P06426
 F-MAMMA1001008//PROGASTRINSIN PRECURSOR (EC 3.4.2
 3.3) (PEPSINOGEN C) (FRAGMENT).//3.2e-14:131:35//M

2941

ACACA FUSCATA FUSCATA (JAPANESE MACAQUE). //P03955
 F-MAMMA1001021//HYPOTHETICAL 35.5 KD PROTEIN IN TR
 ANSPOSON TN4556. //0.016: 61: 42//STREPTOMYCES FRADIA
 E. //P20186
 F-MAMMA1001024
 F-MAMMA1001030//LUTROPIN-CHORIOGONADOTROPIC HORMON
 E RECEPTOR (LH/CG-R) (LSH-R) (LUTEINIZING HORMONE
 RECEPTOR) (FRAGMENT). //2.4e-20: 234: 29//GALLUS GAL
 LUS (CHICKEN). //Q90674
 F-MAMMA1001035//!!!! ALU SUBFAMILY SP WARNING ENTR 10
 Y !!!!!/1.7e-15: 52: 78//HOMO SAPIENS (HUMAN). //P391
 93
 F-MAMMA1001038//NEUROTOXIN II (TOXIN RP-II) (SODIU
 M CHANNEL TOXIN II). //0.53: 25: 48//RADIANTHUS PAUMO
 TENSIS (SEA ANEMONE) (HETERACTIS PAUMOTENSIS). //PO
 1534
 F-MAMMA1001041//SPECTRIN BETA CHAIN, ERYTHROCYTE. /
 /6.3e-18: 112: 43//MUS MUSCULUS (MOUSE). //P15508
 F-MAMMA1001050
 F-MAMMA1001059//PUTATIVE ATP-DEPENDENT RNA HELICAS 20
 E C12C2.06. //1.3e-34: 187: 47//SCHIZOSACCHAROMYCES P
 OMBE (FISSION YEAST). //Q09747
 F-MAMMA1001067//PROTEIN Q300. //0.36: 12: 75//MUS MUS
 CULUS (MOUSE). //Q02722
 F-MAMMA1001073//HEPATOCYTE NUCLEAR FACTOR 3 FORKHE
 AD HOMOLOG 1 (HFB-1). //1.0: 70: 37//RATTUS NORVEGICU
 S (RAT). //Q63244
 F-MAMMA1001074//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //0.00067: 163: 32//HOMO SAPIENS (HUMAN). //P08547
 F-MAMMA1001075//RETINOBLASTOMA BINDING PROTEIN 1 30
 (RBBP-1). //0.53: 72: 34//HOMO SAPIENS (HUMAN). //P293
 74
 F-MAMMA1001078//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //5.0e-79: 184: 73//HOMO SAPIENS (HUMAN). //P08547
 F-MAMMA1001080//IG HEAVY CHAIN PRECURSOR V-III REG
 ION (VH26). //1.7e-27: 82: 71//HOMO SAPIENS (HUMAN). /
 /P01764
 F-MAMMA1001082
 F-MAMMA1001091//HYPOTHETICAL BHLF1 PROTEIN. //3.1e-
 05: 198: 32//EPSTEIN-BARR VIRUS (STRAIN B95-8) (HUMA 40
 N HERPESVIRUS 4). //P03181
 F-MAMMA1001092//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //7.1e-21: 65: 72//HOMO SAPIENS (HUMAN). //P08547
 F-MAMMA1001105//OVO PROTEIN (SHAVEN BABY PROTEIN).
 //1.0e-18: 68: 48//DROSOPHILA MELANOGASTER (FRUIT FL
 Y). //P51521
 F-MAMMA1001110//PROCOLLAGEN ALPHA 1(IV) CHAIN PREC
 URSOR. //0.080: 108: 37//MUS MUSCULUS (MOUSE). //P0246
 3
 F-MAMMA1001126//!!!! ALU SUBFAMILY SB WARNING ENTR 50

2942

Y !!!!!/2.3e-07: 66: 45//HOMO SAPIENS (HUMAN). //P391
 89
 F-MAMMA1001133//HYPOTHETICAL 13.2 KD PROTEIN IN RP
 S4A-BAT2 INTERGENIC REGION. //0.96: 43: 25//SACCHAROM
 YCES CEREVISIAE (BAKER'S YEAST). //P47174
 F-MAMMA1001139//HYPOTHETICAL 36.7 KD PROTEIN AH6.2
 IN CHROMOSOME II. //5.4e-42: 81: 62//CAENORHABDITIS
 ELEGANS. //Q09201
 F-MAMMA1001143//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/0.00014: 36: 66//HOMO SAPIENS (HUMAN). //P3918
 8
 F-MAMMA1001145
 F-MAMMA1001154//CSBA PROTEIN. //1.0: 39: 38//BACILLUS
 SUBTILIS. //P37953
 F-MAMMA1001161//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/5.2e-23: 53: 64//HOMO SAPIENS (HUMAN). //P3918
 8
 F-MAMMA1001162//CD27L RECEPTOR PRECURSOR (T-CELL A
 CTIVATION ANTIGEN CD27). //0.69: 86: 31//MUS MUSCULUS
 (MOUSE). //P41272
 F-MAMMA1001181//HYPOTHETICAL 81.0 KD PROTEIN C35D1
 0.4 IN CHROMOSOME III. //0.00010: 74: 47//CAENORHABDI
 TIS ELEGANS. //Q18486
 F-MAMMA1001186//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!/9.0e-32: 44: 86//HOMO SAPIENS (HUMAN). //P391
 94
 F-MAMMA1001191//OCTAMER-BINDING TRANSCRIPTION FACT
 OR 1 (OTF-1) (NF-A1) (FRAGMENT). //0.096: 40: 40//MAC
 ROPUS EUGENII (TAMMAR WALLABY). //Q28466
 F-MAMMA1001198//EPIDERMAL GROWTH FACTOR RECEPTOR S
 UBSTRATE SUBSTRATE 15 (PROTEIN EPS15) (AF-1P PROTEI
 N). //2.5e-75: 204: 70//HOMO SAPIENS (HUMAN). //P42566
 F-MAMMA1001202//METALLOTHIONEIN-II (MT-II) (MT-IIB
 /MT-IIA). //0.52: 46: 32//CALLINECTES SAPIDUS (BLUE C
 RAB). //P55950
 F-MAMMA1001203//!!!! ALU SUBFAMILY SC WARNING ENTR
 Y !!!!!/7.3e-11: 82: 58//HOMO SAPIENS (HUMAN). //P391
 92
 F-MAMMA1001206//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/1.9e-17: 67: 71//HOMO SAPIENS (HUMAN). //P3918
 8
 F-MAMMA1001215//9 KD PROTEIN. //1.0: 51: 33//HOMO SAP
 IENS (HUMAN). //P13994
 F-MAMMA1001220//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/3.4e-37: 55: 87//HOMO SAPIENS (HUMAN). //P391
 89
 F-MAMMA1001222//HYPOTHETICAL 73.6 KD PROTEIN CY49.
 21. //3.7e-06: 168: 38//MYCOBACTERIUM TUBERCULOSIS. //
 Q10690
 F-MAMMA1001243

2943

2944

- F-MAMMA1001244//TRP OPERON LEADER PEPTIDE.//1.0:1
8:55//SERRATIA MARCESCENS.//P03055
- F-MAMMA1001249//HYPOTHETICAL 7.2 KD PROTEIN IN RPS
2 3' REGION (ORF57).//0.57:23:34//ASTASIA LONGA (EU
GLENOPHYCEAN ALGA).//P34774
- F-MAMMA1001256//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//2.3e-07:79:44//HOMO SAPIENS (HUMAN).//P3918
8
- F-MAMMA1001259//PUTATIVE DNA HELICASE II HOMOLOG
(EC 3.6.1.-).//0.046:86:32//MYCOPLASMA GENITALIUM. 10
//P47486
- F-MAMMA1001260//MYOSIN HEAVY CHAIN, PERINATAL SKEL
ETAL MUSCLE.//2.7e-05:219:27//HOMO SAPIENS (HUMA
N).//P13535
- F-MAMMA1001268//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//9.7e-27:89:67//HOMO SAPIENS (HUMAN).//P08547
- F-MAMMA1001271//ATROPHIN-1 (DENTATORUBRAL-PALLIDOL
UYSIAN ATROPHY PROTEIN).//4.0e-06:126:38//HOMO SAP
IENS (HUMAN).//P54259
- F-MAMMA1001274//!!!! ALU SUBFAMILY SQ WARNING ENTR 20
Y !!!!//7.4e-29:57:66//HOMO SAPIENS (HUMAN).//P391
94
- F-MAMMA1001280//BACTERIOCIN MICROGIN B17 PRECURSOR
(MCB17).//0.27:24:54//ESCHERICHIA COLI.//P05834
- F-MAMMA1001292//HYPOTHETICAL PROTEIN KIAA0176 (FRA
GMENT).//1.3e-73:208:69//HOMO SAPIENS (HUMAN).//Q1
4681
- F-MAMMA1001296//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!//6.9e-22:41:80//HOMO SAPIENS (HUMAN).//P391
93
- F-MAMMA1001298//HYPOTHETICAL PROTEIN HI0371.//0.9
9:29:37//HAEMOPHILUS INFLUENZAE.//P44668
- F-MAMMA1001305//GTPASE-ACTIVATING PROTEIN RHOGAP
(RHO-RELATED SMALL GTPASE PROTEIN ACTIVATOR) (CDC4
2 GTPASE-ACTIVATING PROTEIN) (P50-RHOGAP).//9.9e-6
2:222:54//HOMO SAPIENS (HUMAN).//Q07960
- F-MAMMA1001322//B-CELL GROWTH FACTOR PRECURSOR (BC
GF-12 KD).//2.1e-09:46:60//HOMO SAPIENS (HUMAN).//
P20931
- F-MAMMA1001324//POL POLYPROTEIN [CONTAINS: PROTEAS 40
E (EC 3.4.23.-); REVERSE TRANSCRIPTASE (EC 2.7.7.4
9); RIBONUCLEASE H (EC 3.1.26.4)].//2.5e-43:128:50
//FRIEND MURINE LEUKEMIA VIRUS (ISOLATE PVC-211)
(F-MULV).//P26808
- F-MAMMA1001330//HEMOGLOBIN ZETA CHAIN (FRAGMENTS).
//0.30:51:37//MACROPUSEUGENII (TAMMAR WALLABY).//P
81044
- F-MAMMA1001341//TRISTETRAPROLINE (TTP) (TIS11A) (T
IS11) (ZFP-36) (GROWTHFACTOR- INDUCIBLE NUCLEAR PR
OTEIN NUP475).//0.024:89:39//HOMO SAPIENS (HUMAN). 50
- //P26651
- F-MAMMA1001343//PROBABLE E5 PROTEIN.//0.60:64:29//
HUMAN PAPILOMAVIRUS TYPE 16.//P06927
- F-MAMMA1001346//PROTEINASE INHIBITOR IIB (FRAGMENT
S).//0.97:33:45//SOLANUM TUBEROSUM (POTATO).//P010
82
- F-MAMMA1001383//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!//1.2e-30:86:77//HOMO SAPIENS (HUMAN).//P391
94
- F-MAMMA1001388//LEUCINE-RICH ALPHA-2-GLYCOPROTEIN
(LRG).//9.2e-91:195:92//HOMO SAPIENS (HUMAN).//P02
750
- F-MAMMA1001397//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//3.5e-19:55:69//HOMO SAPIENS (HUMAN).//P3918
8
- F-MAMMA1001408//SALIVARY GLUE PROTEIN SCS-7 PRECUR
SOR.//0.60:45:35//DROSOPHILA MELANOGASTER (FRUIT F
LY).//P02841
- F-MAMMA1001411//GLUCOAMYLASE S1/S2 PRECURSOR (EC
3.2.1.3) (GLUCAN 1,4-ALPHA- GLUCOSIDASE) (1,4-ALPH
A-D-GLUCAN GLUCOHYDROLASE).//5.8e-06:153:26//SACCH
AROMYCES CEREVISIAE (BAKER'S YEAST).//P08640
- F-MAMMA1001419//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!//2.3e-16:99:51//HOMO SAPIENS (HUMAN).//P391
94
- F-MAMMA1001420//!!!! ALU SUBFAMILY SB1 WARNING ENT
RY !!!!//0.0018:23:65//HOMO SAPIENS (HUMAN).//P391
90
- F-MAMMA1001435//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!//1.7e-22:60:58//HOMO SAPIENS (HUMAN).//P391
95
- F-MAMMA1001442
- F-MAMMA1001446//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!//1.2e-23:48:75//HOMO SAPIENS (HUMAN).//P391
94
- F-MAMMA1001452//GENE 35 PROTEIN (GP35).//0.61:31:4
5//MYCOBACTERIOPHAGE L5.//Q05245
- F-MAMMA1001465//HYPOTHETICAL PROTEIN E-115.//0.002
6:68:38//HUMAN ADENOVIRUS TYPE 2.//P03290
- F-MAMMA1001476//URIDINE KINASE (EC 2.7.1.48) (URID
INE MONOPHOSPHOKINASE) (FRAGMENT).//3.7e-94:201:92/
/MUS MUSCULUS (MOUSE).//P52623
- F-MAMMA1001487//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//4.6e-16:89:41//NYCTICEBUS COUCANG (SLOW LORI
S).//P08548
- F-MAMMA1001501//CALPAIN 1, LARGE [CATALYTIC] SUBUN
IT (EC 3.4.22.17) (CALCIUM-ACTIVATED NEUTRAL PROTE
INASE) (CANP) (MU-TYPE).//6.2e-59:86:97//HOMO SAPI
ENS (HUMAN).//P07384
- F-MAMMA1001502//HYPOTHETICAL 11.4 KD PROTEIN (ORF

2945

1). //0.21:79:30//STREPTOMYCES FRADIAE. //P26800
 F-MAMMA1001510
 F-MAMMA1001522//HYPOTHETICAL 35.5 KD PROTEIN IN TR
 ANSPOSON TN4556. //0.67:98:31//STREPTOMYCES FRADIA
 E. //P20186
 F-MAMMA1001547//PROBABLE MOLYBDENUM-PTERIN BINDING
 PROTEIN. //0.97:35:42//HAEMOPHILUS INFLUENZAE. //P4
 5183
 F-MAMMA1001551//HYPOTHETICAL PROTEIN MJ0458.1. //0.
 038:31:41//METHANOCOCCUS JANNASCHII. //P81308
 F-MAMMA1001575
 F-MAMMA1001576//TUBULIN GAMMA CHAIN. //1.6e-86:162:
 99//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P23330
 F-MAMMA1001590//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!//0.0035:38:55//HOMO SAPIENS (HUMAN). //P3919
 5
 F-MAMMA1001600//CONNECTIVE TISSUE GROWTH FACTOR PR
 ECURSOR. //0.85:53:33//HOMO SAPIENS (HUMAN). //P2927
 9
 F-MAMMA1001604//HYPOTHETICAL 11.1 KD PROTEIN C3OD1
 1.02C IN CHROMOSOME I. //0.14:82:29//SCHIZOSACCHARO
 MYCES POMBE (FISSION YEAST). //Q09902
 F-MAMMA1001606//HIGH MOBILITY GROUP PROTEIN HMGI-
 C. //8.2e-05:77:37//HOMOSAPIENS (HUMAN). //P52926
 F-MAMMA1001620//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!//5.5e-05:24:66//HOMO SAPIENS (HUMAN). //P391
 95
 F-MAMMA1001627//CUTICLE COLLAGEN 40. //0.82:131:31/
 /CAENORHABDITIS ELEGANS. //P34804
 F-MAMMA1001630//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//8.6e-26:57:78//HOMO SAPIENS (HUMAN). //P391
 94
 F-MAMMA1001633//ZINC FINGER PROTEIN 165. //6.9e-38:
 160:55//HOMO SAPIENS (HUMAN). //P49910
 F-MAMMA1001635
 F-MAMMA1001649//SPERM PROTAMINE P1. //0.39:31:41//T
 ACHYGLOSSUS ACULEATUSACULEATUS (AUSTRALIAN ECHIDN
 A). //P35311
 F-MAMMA1001654//NON-RECEPTOR TYROSINE KINASE SPORE
 LYSIS A (EC 2.7.1.112) (TYROSINE- PROTEIN KINASE
 1). //5.6e-06:99:28//DICTYOSTELIUM DISCOIDEUM(SLIME
 MOLD). //P18160
 F-MAMMA1001663//VERY HYPOTHETICAL XYLU PROTEIN. //
 0.99:27:37//ESCHERICHIACOLI. //P05056
 F-MAMMA1001670//CUTICLE COLLAGEN 1. //0.033:97:37//
 CAENORHABDITIS ELEGANS. //P08124
 F-MAMMA1001671
 F-MAMMA1001679//PROCOLLAGEN ALPHA 2(IV) CHAIN PREC
 URSOR. //0.92:32:50//HOMO SAPIENS (HUMAN). //P08572
 F-MAMMA1001683//HYPOTHETICAL 35.5 KD PROTEIN IN TR

2946

ANSPOSON TN4556. //0.00026:147:34//STREPTOMYCES FRA
 DIAE. //P20186
 F-MAMMA1001686
 F-MAMMA1001692//SMALL HYDROPHOBIC PROTEIN (SMALL P
 ROTEIN 1A). //1.0:34:26//BOVINE RESPIRATORY SYNCYTI
 AL VIRUS (STRAIN A51908) (BRS). //P24616
 F-MAMMA1001711//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//1.1e-28:56:69//HOMO SAPIENS (HUMAN). //P391
 94
 10 F-MAMMA1001715//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!//1.6e-08:39:71//HOMO SAPIENS (HUMAN). //P3918
 8
 F-MAMMA1001730//METALLOTHIONEIN-B (MTB). //1.0:17:6
 4//STRONGYLOCENTROTUSPURPURATUS (PURPLE SEA URCHI
 N). //Q27287
 F-MAMMA1001735//TUBULIN BETA-5 CHAIN (CLASS-V). //
 5.1e-121:213:97//GALLUSGALLUS (CHICKEN). //P09653
 F-MAMMA1001740
 F-MAMMA1001743//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!//1.3e-09:100:42//HOMO SAPIENS (HUMAN). //P39
 195
 F-MAMMA1001744//POU DOMAIN PROTEIN 2. //0.97:59:38/
 /BRACHYDANIO RERIO (ZEBRAFISH) (ZEBRA DANI0). //Q90
 270
 F-MAMMA1001745//LINE-1 REVERSE TRANSCRIPTASE HOMOL
 OG. //2.1e-43:199:42//HOMO SAPIENS (HUMAN). //P08547
 F-MAMMA1001751//TWK-8 PROTEIN. //2.9e-15:77:36//CAE
 NORHABDITIS ELEGANS. //P34410
 F-MAMMA1001754//MALE SPECIFIC SPERM PROTEIN MST84D
 D. //0.019:20:45//DROSOPHILA MELANOGASTER (FRUIT FL
 Y). //Q01645
 F-MAMMA1001757//HYPOTHETICAL 9.2 KD PROTEIN IN RNP
 A 3' REGION. //0.94:30:43//PSEUDOMONAS PUTIDA. //P257
 53
 F-MAMMA1001760//!!!! ALU SUBFAMILY SB2 WARNING ENT
 RY !!!!!//4.6e-34:103:59//HOMO SAPIENS (HUMAN). //P3
 9191
 F-MAMMA1001764
 F-MAMMA1001768//HYPOTHETICAL PROTEIN UL61. //0.042:
 167:33//HUMAN CYTOMEGALOVIRUS (STRAIN AD169). //P16
 818
 F-MAMMA1001769//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//2.0e-29:97:69//HOMO SAPIENS (HUMAN). //P391
 94
 F-MAMMA1001771//TRANSMEMBRANE PROTEIN SEX PRECURSO
 R. //3.3e-09:123:32//HOMO SAPIENS (HUMAN). //P51805
 F-MAMMA1001783//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!//1.1e-09:55:61//HOMO SAPIENS (HUMAN). //P3918
 8
 50 F-MAMMA1001785//RAS-RELATED PROTEIN RABC. //1.9e-0

2947

2948

6: 120: 25//DICTYOSTELIUMDISCOIDEUM (SLIME MOLD). //P
34143

F-MAMMA1001788//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG. //9. 3e-29: 46: 76//HOMO SAPIENS (HUMAN). //P08547

F-MAMMA1001790//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//4. 7e-24: 69: 69//HOMO SAPIENS (HUMAN). //P3918

8

F-MAMMA1001806//HYPOTHETICAL 21.2 KD PROTEIN IN TO
R2-MNN4 INTERGENIC REGION. //0. 95: 58: 36//SACCHAROMY
CES CEREVISIAE (BAKER'S YEAST). //P36042

F-MAMMA1001812//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!//8. 8e-12: 53: 69//HOMO SAPIENS (HUMAN). //P391

95

F-MAMMA1001815//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG. //0. 11: 30: 70//HOMOSAPIENS (HUMAN). //P08547

F-MAMMA1001817//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//2. 9e-16: 86: 55//HOMO SAPIENS (HUMAN). //P3918

8

F-MAMMA1001818

F-MAMMA1001820//VITELLINE MEMBRANE PROTEIN VM26AB
PRECURSOR (PROTEIN TU-4) (PROTEIN SV23). //0. 0030: 6
3: 42//DROSOPHILA MELANOGASTER (FRUIT FLY). //P13238

F-MAMMA1001824//APTOTOXIN VII (PARALYTIC PEPTIDE V
II) (PP VII). //0. 99: 26: 34//APTOSTICHUS SCHLINGERI
(TRAP-DOOR SPIDER). //P49271

F-MAMMA1001836//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!//2. 6e-35: 77: 88//HOMO SAPIENS (HUMAN). //P391

95

F-MAMMA1001837//ZINC FINGER PROTEIN 191. //1. 3e-27:
106: 58//HOMO SAPIENS (HUMAN). //014754

F-MAMMA1001848//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//1. 0e-19: 92: 58//HOMO SAPIENS (HUMAN). //P3918

8

F-MAMMA1001851

F-MAMMA1001854

F-MAMMA1001858//ISOTOCIN-NEUROPHYSIN IT 1 PRECURSO
R. //0. 93: 42: 38//CATOSTOMUS COMMERSONI (WHITE SUCKE
R). //P15210

F-MAMMA1001864//PROBABLE ABC TRANSPORTER PERMEASE
PROTEIN MG189. //0. 77: 161: 27//MYCOPLASMA GENITALIU
M. //P47435

F-MAMMA1001868//FK506-BINDING NUCLEAR PROTEIN (PEP
TIDYL-PROLYL CIS-TRANSISOMERASE) (PPIASE) (EC 5. 2.
1. 8) (PROLINE ROTAMASE) (NUCLEOLAR PROLINEISOMERAS
E) (FKBP-70). //0. 00013: 219: 26//SACCHAROMYCES CEREV
ISIAE (BAKER'S YEAST). //P38911

F-MAMMA1001874//SPERM HISTONE P2 PRECURSOR (PROTAM
INE MP2). //0. 0075: 76: 31//MUS MUSCULUS (MOUSE). //PO
7978

F-MAMMA1001878//GLYCINE-RICH CELL WALL STRUCTURAL 50

PROTEIN (CLONE W10-1) (FRAGMENT). //0. 020: 10: 80//LYC
OPERSICON ESCULENTUM (TOMATO). //Q01157

F-MAMMA1001880

F-MAMMA1001890//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!//5. 1e-34: 56: 83//HOMO SAPIENS (HUMAN). //P391
92

F-MAMMA1001907//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//2. 7e-12: 44: 68//HOMO SAPIENS (HUMAN). //P391
94

10 F-MAMMA1001908//HYPOTHETICAL 16.2 KD PROTEIN IN PR
P24-RRN9 INTERGENIC REGION. //0. 00013: 77: 37//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST). //Q03525

F-MAMMA1001931//HYPOTHETICAL 118.2 KD PROTEIN F43C
1.1 IN CHROMOSOME III. //0. 41: 106: 29//CAENORHABDITI
S ELEGANS. //Q09564

F-MAMMA1001956//OCTAPEPTIDE-REPEAT PROTEIN T2. //0.
00053: 149: 30//MUS MUSCULUS (MOUSE). //Q06666

F-MAMMA1001963//HYPOTHETICAL PROTEIN IN NAC 5' REGI
ON (ORF X) (FRAGMENT). //1. 0: 46: 28//KLEBSIELLA AERO
GENES. //Q08600

F-MAMMA1001969//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG. //1. 7e-34: 97: 68//HOMO SAPIENS (HUMAN). //P08547

F-MAMMA1001970//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG. //6. 2e-07: 67: 37//HOMO SAPIENS (HUMAN). //P08547

F-MAMMA1001992//PROTEIN Q300. //0. 53: 14: 71//MUS MUS
CULUS (MOUSE). //Q02722

F-MAMMA1002009//PROBABLE E5 PROTEIN. //0. 17: 56: 32//
HUMAN PAPILLOMAVIRUS TYPE 31. //P17385

F-MAMMA1002011//MYRISTOYLATED ALANINE-RICH C-KINAS
E SUBSTRATE (MARCKS) (PROTEIN KINASE C SUBSTRATE,
80 KD PROTEIN, LIGHT CHAIN) (PKCSL) (80K-L PROTEI
N). //1. 0: 100: 31//HOMO SAPIENS (HUMAN). //P29966

30

F-MAMMA1002032//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//3. 1e-21: 86: 65//HOMO SAPIENS (HUMAN). //P3918

8

F-MAMMA1002033//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//8. 5e-20: 67: 58//HOMO SAPIENS (HUMAN). //P3918

8

F-MAMMA1002041//MALE SPECIFIC SPERM PROTEIN MST84D
C. //1. 0: 17: 52//DROSOPHILA MELANOGASTER (FRUIT FL
Y). //Q01644

F-MAMMA1002042//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!!//0. 19: 45: 46//HOMO SAPIENS (HUMAN). //P39192

F-MAMMA1002047//TYROSINE AMINOTRANSFERASE (EC 2. 6.
1. 5) (L-TYROSINE: 2-OXOGLUTARATE AMINOTRANSFERASE)
(TAT). //0. 0017: 50: 46//RATTUS NORVEGICUS (RAT). //PO
4694

F-MAMMA1002056//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//3. 2e-37: 70: 77//HOMO SAPIENS (HUMAN). //P391

94

94

2949

F-MAMMA1002058//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//1.1e-08:26:76//HOMO SAPIENS (HUMAN).//P3918
8

F-MAMMA1002068//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//2.0e-11:78:46//HOMO SAPIENS (HUMAN).//P08547
F-MAMMA1002078//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//0.96:26:46//COTURNIX COTURNIX JAPONICA
(JAPANESE QUAIL).//P50682

F-MAMMA1002082//SUPPRESSOR PROTEIN SRP40.//0.23:9
5:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P
32583

F-MAMMA1002084//HYPOTHETICAL 7.5 KD PROTEIN.//1.0:
40:35//VACCINIA VIRUS(STRAIN COPENHAGEN).//P20520

F-MAMMA1002093

F-MAMMA1002108//HYPOTHETICAL 35.5 KD PROTEIN IN TR
ANSPOSON TN4556.//0.00079:143:33//STREPTOMYCES FRA
DIAE.//P20186

F-MAMMA1002118//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//1.0:43:34//METRIDIVM SENILE (BROWN SEA
ANEMONE) (FRILLED SEA ANEMONE).//047493

F-MAMMA1002125//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!//1.9e-14:60:68//HOMO SAPIENS (HUMAN).//P391
92

F-MAMMA1002132

F-MAMMA1002140//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//1.4e-24:69:65//HOMO SAPIENS (HUMAN).//P3918
8

F-MAMMA1002143//SERUM PROTEIN MSE55.//2.1e-16:166:
43//HOMO SAPIENS (HUMAN).//Q00587

F-MAMMA1002145//36.4 KD PROLINE-RICH PROTEIN.//0.0
0014:84:29//LYCOPERSICON ESCULENTUM (TOMATO).//Q00
451

F-MAMMA1002153

F-MAMMA1002155

F-MAMMA1002156//METALLOPROTEINASE INHIBITOR PRECUR
SOR.//0.90:58:34//STREPTOMYCES NIGRESCENS.//P01077

F-MAMMA1002158

F-MAMMA1002170//40S RIBOSOMAL PROTEIN S2 (S4) (LLR
EP3 PROTEIN).//6.0e-66:157:70//HOMO SAPIENS (HUMA
N).//P15880

F-MAMMA1002174//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//6.5e-25:56:64//HOMO SAPIENS (HUMAN).//P3918
8

F-MAMMA1002198//THIOREDOXIN PEROXIDASE 1 (THIOREDO
XIN-DEPENDENT PEROXIDEREDUCTASE 1) (THIOL-SPECIFIC
ANTIOXIDANT PROTEIN) (TSA) (PRP) (NATURALKILLER C
ELL ENHANCING FACTOR B) (NKEF-B).//9.0e-09:28:100/
/HOMO SAPIENS(HUMAN).//P32119

F-MAMMA1002209//TRANSCRIPTION INITIATION FACTOR TF
IID 135 KD SUBUNIT (TAFII-135) (TAFII135) (TAFII-1

2950

30) (TAFII130).//0.0023:132:33//HOMO SAPIENS(HUMA
N).//000268

F-MAMMA1002215//PROCOLLAGEN ALPHA 1(I) CHAIN PRECU
RSOR.//0.00032:68:35//HOMO SAPIENS (HUMAN).//P0245
2

F-MAMMA1002219//INTRACELLULAR PROTEIN TRANSPORT PR
OTEIN US01.//0.0079:224:24//SACCHAROMYCES CEREVISI
AE (BAKER'S YEAST).//P25386

F-MAMMA1002230

10 F-MAMMA1002236//TRANSLATION INITIATION FACTOR EIF-
2B GAMMA SUBUNIT (EIF-2B GDP-GTP EXCHANGE FACTOR).
//1.4e-118:151:94//RATTUS NORVEGICUS (RAT).//P7054
1

F-MAMMA1002243//WISKOTT-ALDRICH SYNDROME PROTEIN H
OMOLOG (WASP).//0.028:112:33//MUS MUSCULUS (MOUS
E).//P70315

F-MAMMA1002250//T-CELL RECEPTOR BETA CHAIN PRECURS
OR (ANA 11).//0.0012:80:32//ORYCTOLAGUS CUNICULUS
(RABBIT).//P06333

20 F-MAMMA1002267//ATP SYNTHASE A CHAIN (EC 3.6.1.34)
(PROTEIN 6).//0.17:139:28//TRYPANOSOMA BRUCEI BRU
CEI.//P24499

F-MAMMA1002268//60S RIBOSOMAL PROTEIN L22.//0.0002
6:163:30//DROSOPHILA MELANOGASTER (FRUIT FLY).//P5
0887

F-MAMMA1002269//HISTIDINE-RICH, METAL BINDING POLY
PEPTIDE.//0.35:14:57//HELICOBACTER PYLORI (CAMPYLO
BACTER PYLORI).//Q48251

30 F-MAMMA1002282//!!!! ALU SUBFAMILY SC WARNING ENTR
Y !!!!//6.1e-05:32:65//HOMO SAPIENS (HUMAN).//P391
92

F-MAMMA1002292//TROPOMYOSIN 2.//1.4e-05:100:30//SA
CCHAROMYCES CEREVISIAE(BAKER'S YEAST).//P40414

F-MAMMA1002293//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//6.8e-25:127:44//HOMO SAPIENS (HUMAN).//P391
88

F-MAMMA1002294//ALPHA TRANS-INDUCING PROTEIN (ALPH
A-TIF).//0.00011:138:38//BOVINE HERPESVIRUS TYPE 1
(STRAIN P8-2).//P30020

40 F-MAMMA1002297//A-AGGLUTININ ATTACHMENT SUBUNIT PR
ECURSOR.//0.15:144:30//SACCHAROMYCES CEREVISIAE (B
AKER'S YEAST).//P32323

F-MAMMA1002298//PROLINE-RICH PROTEIN MP-3 (FRAGEN
T).//1.0e-05:40:50//MUS MUSCULUS (MOUSE).//P05143

F-MAMMA1002299//NADH-UBIQUINONE OXIDOREDUCTASE CHA
IN 4L (EC 1.6.5.3).//0.84:65:32//STRUTHIO CAMELUS
(OSTRICH).//021405

F-MAMMA1002308//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//4.3e-29:61:73//HOMO SAPIENS (HUMAN).//P3918

50 8

2951

F-MAMMA1002310//SPERM MITOCHONDRIAL CAPSULE SELENO
PROTEIN (MCS).//0.00016:70:38//MUS MUSCULUS (MOUS
E).//P15265

F-MAMMA1002311//!!!! ALU SUBFAMILY SB WARNING ENTR
Y !!!!!/9.4e-09:84:54//HOMO SAPIENS (HUMAN).//P391
89

F-MAMMA1002312//HYPOTHETICAL 10.8 KD PROTEIN IN CP
30-RIII INTERGENIC REGION (URF Y).//0.48:48:33//BA
CTERIOPHAGE T4.//P33084

F-MAMMA1002317

F-MAMMA1002319//RETROVIRUS-RELATED POL POLYPROTEIN
[CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); E
NDONUCLEASE].//0.011:128:27//MUS MUSCULUS (MOUSE).
//P11369

F-MAMMA1002322//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/5.2e-20:92:57//HOMO SAPIENS (HUMAN).//P391
95

F-MAMMA1002329//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//0.051:33:36//XENOPUS LAEVIS (AFRICAN CL
AWED FROG).//P03931

F-MAMMA1002332//LINE-1 REVERSE TRANSCRIPTASE HOMOL
OG.//6.5e-20:116:51//HOMO SAPIENS (HUMAN).//P08547

F-MAMMA1002333//COLLAGEN ALPHA 1(I) CHAIN (FRAGMEN
TS).//0.0017:214:31//BOS TAURUS (BOVINE).//P02453

F-MAMMA1002339//COPPER-METALLOTHIONEIN (CU-MT).//
0.59:42:38//HELIIX POMATIA (ROMAN SNAIL) (EDIBLE SN
AIL).//P55947

F-MAMMA1002347//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/0.43:26:61//HOMO SAPIENS (HUMAN).//P39188

F-MAMMA1002351//HYPOTHETICAL PROTEIN MJ0304.//2.3e
-07:139:25//METHANOCOCCUS JANNASCHII.//Q57752

F-MAMMA1002352

F-MAMMA1002353//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/0.00028:31:80//HOMO SAPIENS (HUMAN).//P3918
8

F-MAMMA1002355//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/4.2e-28:87:73//HOMO SAPIENS (HUMAN).//P391
93

F-MAMMA1002356//RELAXIN.//0.95:31:35//SQUALUS ACAN
THIAS (SPINY DOGFISH).//P11953

F-MAMMA1002359//CHLOROPLAST 50S RIBOSOMAL PROTEIN
L33.//0.93:44:36//GUILLARDIA THETA (CRYPTOMONAS PH
I).//078487

F-MAMMA1002360//LATE L2 MU CORE PROTEIN PRECURSOR
(PROTEIN X).//0.94:30:43//BOVINE ADENOVIRUS TYPE 2
(MASTADENOVIRUS BOS2).//Q96626

F-MAMMA1002361//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/2.0e-08:45:68//HOMO SAPIENS (HUMAN).//P3918
8

F-MAMMA1002362//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3 50

2952

4) (A6L).//0.58:23:26//LUMBRICUS TERRESTRIS (COMMO
N EARTHWORM).//Q34942

F-MAMMA1002380//SALIVARY GLUE PROTEIN SGS-3 PRECUR
SOR.//0.23:100:27//DROSOPHILA SIMULANS (FRUIT FL
Y).//P13729

F-MAMMA1002384

F-MAMMA1002385//HYPOTHETICAL 40.9 KD PROTEIN IN OR
C2-TIP1 INTERGENIC REGION.//3.8e-14:125:37//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P38241

10 F-MAMMA1002392//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//1.0:17:58//BRANCHIOSTOMA LANCEOLATUM (C
OMMON LANCELET) (AMPHIOXUS).//021003

F-MAMMA1002411//30S RIBOSOMAL PROTEIN S17.//0.85:4
9:32//SYNECHOCYSTIS SP. (STRAIN PCC 6803).//P73311

F-MAMMA1002413//NADH-UBIQUINONE OXIDOREDUCTASE CHA
IN 1 (EC 1.6.5.3) (FRAGMENT).//0.97:41:39//DROSOPH
ILA AFFINIS (FRUIT FLY).//P51926

F-MAMMA1002417//RFBJ PROTEIN.//0.99:31:35//SHIGELL
A FLEXNERI.//P37786

20 F-MAMMA1002427//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/1.6e-33:135:59//HOMO SAPIENS (HUMAN).//P39
194

F-MAMMA1002428//HYPOTHETICAL PROTEIN C18.//0.97:3
4:44//SWINEPOX VIRUS (STRAIN KASZA) (SPV).//P32217

F-MAMMA1002434//!!!! ALU SUBFAMILY SB WARNING ENTR
Y !!!!!/3.1e-36:56:78//HOMO SAPIENS (HUMAN).//P391
89

F-MAMMA1002446

F-MAMMA1002454//EARLY NODULIN 20 PRECURSOR (N-20).
//0.77:57:45//MEDICAGOTRUNCATULA (BARREL MEDIC).//
P93329

F-MAMMA1002461//VASODILATOR-STIMULATED PHOSPHOPROT
EIN (VASP).//1.3e-05:193:32//CANIS FAMILIARIS (DO
G).//P50551

F-MAMMA1002470//HYPOTHETICAL 80.7 KD PROTEIN IN ER
G7-NMD2 INTERGENIC REGION.//1.0e-75:231:60//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P38795F-MAMMA
1002475//POSSIBLE GLOBAL TRANSCRIPTION ACTIVATOR S

40 NF2L4 (SNF2-BETA) (BRG-1 PROTEIN) (MITOTIC GROWTH
AND TRANSCRIPTION ACTIVATOR) (BRAHMAPROTEIN HOMOLO
G 1).//0.013:99:30//HOMO SAPIENS (HUMAN).//P51532

F-MAMMA1002480//NONSTRUCTURAL PROTEIN 5B.//1.0:23:
43//HUMAN CORONAVIRUS(STRAIN 229E).//P19741

F-MAMMA1002485//STANNIOCALCIN PRECURSOR.//2.1e-23:
88:46//HOMO SAPIENS (HUMAN).//P52823

F-MAMMA1002494//MOLT-INHIBITING HORMONE (MIH).//1.
0:32:37//PROCAMBARUS CLARKII (RED SWAMP CRAYFISH).
//P55848

F-MAMMA1002498//6.7 KD PROTEIN (ORF 5).//1.0:26:42
//BARLEY YELLOW DWARFVIRUS (ISOLATE PAV) (BYDV).//

- P09517
F-MAMMA1002524//HYPOTHETICAL 117.8 KD PROTEIN IN S
TE2-FRS2 INTERGENIC REGION.//5.0e-26:222:35//SACCH
AROMYCES CEREVISIAE (BAKER'S YEAST).//P43571
F-MAMMA1002530//CYTOSOLIC PHOSPHOLIPASE A2 (EC 3.
1.1.4) (CPLA2) (PHOSPHATIDYLCHOLINE 2-ACYLHYDROLAS
E) / LYSOPHOSPHOLIPASE (EC 3.1.1.5).//4.5e-12:88:4
4//HOMO SAPIENS (HUMAN).//P47712
F-MAMMA1002545//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/4.3e-29:97:71//HOMO SAPIENS (HUMAN).//P391 10
95
F-MAMMA1002554//DNA-DIRECTED RNA POLYMERASE II LAR
GEST SUBUNIT (EC 2.7.7.6) (RPB1) (FRAGMENT).//0.4
6:54:40//CRICETULUS GRISEUS (CHINESE HAMSTER).//P1
1414
F-MAMMA1002556//METALLOTHIONEIN 20-I ISOFORMS A AN
D B (MT-20-1A AND MT-20-1B).//0.99:21:47//MYTILUS
EDULIS (BLUE MUSSEL).//P80251
F-MAMMA1002566//TRANSCRIPTION FACTOR P65 (NUCLEAR
FACTOR NF-KAPPA-B P65SUBUNIT).//0.70:130:30//MUS M 20
USCULUS (MOUSE).//Q04207
F-MAMMA1002571//DNA-DIRECTED RNA POLYMERASE II LAR
GEST SUBUNIT (EC 2.7.7.6) (FRAGMENT).//0.54:45:51/
/DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P35084
F-MAMMA1002573//PARATHYMOSIN.//1.5e-07:69:46//HOMO
SAPIENS (HUMAN).//P20962
F-MAMMA1002585//MYOSIN LIGHT CHAIN 1, SLOW-TWITCH
MUSCLE B/VENTRICULAR ISOFORM (FRAGMENT).//0.38:36:
36//MUS MUSCULUS (MOUSE).//P09542
F-MAMMA1002590//!!!! ALU SUBFAMILY SX WARNING ENTR 30
Y !!!!!/0.99:22:77//HOMO SAPIENS (HUMAN).//P39195
F-MAMMA1002597//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/2.1e-18:44:70//HOMO SAPIENS (HUMAN).//P391
94
F-MAMMA1002598//60S RIBOSOMAL PROTEIN L7.//1.8e-1
6:40:100//HOMO SAPIENS (HUMAN).//P18124
F-MAMMA1002603
F-MAMMA1002612//30S RIBOSOMAL PROTEIN S16 (FRAGMEN
T).//1.0:29:37//THERMUS AQUATICUS.//007348
F-MAMMA1002617//ACIDIC PROLINE-RICH PROTEIN PRECUR 40
SOR (CLONE PRP25) (FRAGMENT).//0.00041:81:34//RATT
US NORVEGICUS (RAT).//P10164
F-MAMMA1002618//ESCARGOT/SLAIL PROTEIN HOMOLOG (FR
AGMENT).//0.11:18:50//PSYCHODA CINEREA.//Q02027
F-MAMMA1002619//PROBABLE UBIQUITIN CARBOXYL-TERMIN
AL HYDROLASE K02C4.3 (EC 3.1.2.15) (UBIQUITIN THIO
LESTERASE) (UBIQUITIN-SPECIFIC PROCESSING PROTEAS
E) (DEUBIQUITINATING ENZYME).//1.8e-13:110:40//CAE
NORHABDITIS ELECANS.//Q09931
F-MAMMA1002622//!!!! ALU SUBFAMILY J WARNING ENTRY 50 8
!!!!/8.4e-05:53:58//HOMO SAPIENS (HUMAN).//P3918
8
F-MAMMA1002623//PEPTIDYL-GLYCINE ALPHA-AMIDATING M
ONOOXYGENASE PRECURSOR (EC 1.14.17.3) (PAM).//2.6e-
07:37:78//HOMO SAPIENS (HUMAN).//P19021
F-MAMMA1002625
F-MAMMA1002629//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/4.4e-19:49:73//HOMO SAPIENS (HUMAN).//P3918
8
F-MAMMA1002636//COLLAGEN ALPHA 2(VI) CHAIN (FRAGME
NT).//1.7e-07:189:32//HOMO SAPIENS (HUMAN).//P1211
0
F-MAMMA1002637//KINESIN LIGHT CHAIN (KLC).//7.7e-5
4:227:52//RATTUS NORVEGICUS (RAT).//P37285
F-MAMMA1002646//NEUROFILAMENT TRIPLET H PROTEIN (2
00 KD NEUROFILAMENT PROTEIN) (NF-H).//0.034:199:25
//MUS MUSCULUS (MOUSE).//P19246
F-MAMMA1002650//TRANSCRIPTION REGULATOR PROTEIN BA
CH2 (BTB AND CNC HOMOLOG 2).//1.7e-07:104:32//MUS
MUSCULUS (MOUSE).//P97303
F-MAMMA1002655//SMALL PROLINE-RICH PROTEIN II (SPR
-II) (CLONE 174N).//1.0:25:44//HOMO SAPIENS (HUMA
N).//P22532
F-MAMMA1002662
F-MAMMA1002665//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!/2.3e-07:54:57//HOMO SAPIENS (HUMAN).//P391
94
F-MAMMA1002671//ACETYL-COENZYME A SYNTHETASE (EC
6.2.1.1) (ACETATE--COALICASE) (ACYL- ACTIVATING EN
ZYME).//1.4e-10:144:31//ESCHERICHIA COLI.//P27550
F-MAMMA1002673//BREVICAN CORE PROTEIN PRECURSOR.//
0.76:64:39//BOS TAURUS (BOVINE).//Q28062
F-MAMMA1002684//HYPOTHETICAL 11.8 KD PROTEIN IN GP
55-NRDC INTERGENIC REGION.//0.094:77:27//BACTERIOP
HAGE T4.//P07079
F-MAMMA1002685//COLLAGEN ALPHA 1(I) CHAIN (FRAGMEN
TS).//0.0017:177:34//RATTUS NORVEGICUS (RAT).//P02
454
F-MAMMA1002698
F-MAMMA1002699//HYPOTHETICAL 45.1 KD PROTEIN IN RP
S5-ZMS1 INTERGENIC REGION.//1.2e-28:127:47//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P47160
F-MAMMA1002701//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/1.0:14:92//HOMOSAPIENS (HUMAN).//P39188
F-MAMMA1002708//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/7.9e-27:52:65//HOMO SAPIENS (HUMAN).//P391
93
F-MAMMA1002711//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!/3.7e-24:54:75//HOMO SAPIENS (HUMAN).//P3918

2955

F-MAMMA1002721
 F-MAMMA1002727//SOX-13 PROTEIN (FRAGMENT).//0.70:3
 6:38//MUS MUSCULUS (MOUSE).//Q04891
 F-MAMMA1002728//HYPOTHETICAL 6.0 KD PROTEIN.//1.0:
 25:44//THERMOPROTEUS TENAX VIRUS 1 (STRAIN KRA1)
 (TTV1).//P19305
 F-MAMMA1002744//HYPOTHETICAL 13.4 KD PROTEIN IN AC
 T5-YCK1 INTERGENIC REGION.//1.0:52:34//SACCHAROMYC
 ES CEREVISIAE (BAKER'S YEAST).//P38834
 F-MAMMA1002746//HYPOTHETICAL 5.6 KD PROTEIN (ORF A 10
 -45).//1.0:22:40//SULFOLOBUS VIRUS-LIKE PARTICLE S
 SV1.//P20198
 F-MAMMA1002748
 F-MAMMA1002754//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/5.1e-21:56:64//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1002758//MALE SPECIFIC SPERM PROTEIN MST84D
 D.//0.37:14:64//DROSOPHILA MELANOGASTER (FRUIT FL
 Y).//Q01645
 F-MAMMA1002764//!!!! ALU SUBFAMILY SQ WARNING ENTR 20
 Y !!!!!/4.7e-32:79:60//HOMO SAPIENS (HUMAN).//P391
 94
 F-MAMMA1002765//PARATHYMSIN.//0.79:63:28//BOS TAU
 RUS (BOVINE).//P08814
 F-MAMMA1002769//GAR2 PROTEIN.//0.00037:192:27//SCH
 IZOSACCHAROMYCES POMBE(FISSION YEAST).//P41891
 F-MAMMA1002775//HYPOTHETICAL 36.7 KD PROTEIN C2F7.
 14C IN CHROMOSOME I.//5.4e-54:240:49//SCHIZOSACCHA
 ROMYCES POMBE (FISSION YEAST).//Q09704
 F-MAMMA1002780
 F-MAMMA1002782//MARGATOXIN (MCTX).//1.0:31:38//CEN
 TRUROIDES MARGARITATUS(SCORPION).//P40755
 F-MAMMA1002796//ICE NUCLEATION PROTEIN.//0.0018:10
 0:41//PSEUDOMONAS FLUORESCENS.//P09815
 F-MAMMA1002807//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/9.3e-23:100:59//HOMO SAPIENS (HUMAN).//P391
 88
 F-MAMMA1002820//NEUROTOXIN IV (LQ IV).//1.0:18:50
 //LEIURUS QUINQUESTRIATUS QUINQUESTRIATUS (EGYPTIA
 N SCORPION).//P01489
 F-MAMMA1002830//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!/4.7e-24:55:74//HOMO SAPIENS (HUMAN).//P391
 95
 F-MAMMA1002833//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/2.6e-31:95:73//HOMO SAPIENS (HUMAN).//P391
 89
 F-MAMMA1002835//HYPOTHETICAL 42.1 KD PROTEIN F13G
 3.3 IN CHROMOSOME I.//1.0:54:37//CAENORHABDITIS EL
 EGANS.//Q19417
 F-MAMMA1002838//!!!! ALU SUBFAMILY SP WARNING ENTR 50

2956

Y !!!!!/2.5e-27:99:70//HOMO SAPIENS (HUMAN).//P391
 93
 F-MAMMA1002842//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!/2.3e-13:65:63//HOMO SAPIENS (HUMAN).//P391
 95
 F-MAMMA1002843//METALLOTHIONEIN-II (MT-II).//0.97:
 19:47//MUS MUSCULUS (MOUSE).//P02798
 F-MAMMA1002844//HYPOTHETICAL 24.1 KD PROTEIN IN LE
 F4-P33 INTERGENIC REGION.//4.9e-08:119:36//AUTOGRA
 PHA CALIFORNICA NUCLEAR POLYHEDROSIS VIRUS (ACMNP
 V).//P41479
 F-MAMMA1002858//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
 4) (A6L).//0.98:37:37//PAN TROGLODYTES (CHIMPANZE
 E).//Q35647
 F-MAMMA1002868//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/3.8e-10:51:62//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1002869//PINCH PROTEIN (PARTICULARY INTERES
 TING NEW CYS-HIS PROTEIN).//1.8e-95:194:78//HOMO S
 APIENS (HUMAN).//P48059
 F-MAMMA1002871//G-PROTEIN COUPLED RECEPTOR HOMOLOG
 R33.//1.0:51:35//RATCYTOMEGALOVIRUS (STRAIN MAAST
 RICHT).//012000
 F-MAMMA1002880
 F-MAMMA1002881//GLIOMA PATHOGENESIS-RELATED PROTEI
 N (RTVP-1 PROTEIN).//3.3e-22:180:35//HOMO SAPIENS
 (HUMAN).//P48060
 F-MAMMA1002886//MYOSIN HEAVY CHAIN IB (MYOSIN HEAV
 Y CHAIN IL).//0.00011:148:39//ACANTHAMOEBA CASTELL
 ANII (AMOEBIA).//P19706
 F-MAMMA1002887
 F-MAMMA1002890//A-AGGLUTININ ATTACHMENT SUBUNIT PR
 ECURSOR.//0.030:142:25//SACCHAROMYCES CEREVISIAE
 (BAKER'S YEAST).//P32323
 F-MAMMA1002892
 F-MAMMA1002895//HYPOTHETICAL PROTEIN UL61.//0.0009
 9:143:35//HUMAN CYTOMEGALOVIRUS (STRAIN AD169).//P
 16818
 F-MAMMA1002908//T-CELL RECEPTOR BETA CHAIN PRECURS
 OR (ANA 11).//0.12:44:43//ORYCTOLAGUS CUNICULUS (R
 ABBIT).//P06333
 F-MAMMA1002909//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/0.00011:28:75//HOMO SAPIENS (HUMAN).//P3918
 8
 F-MAMMA1002930//BOMBYXIN A-7 PRECURSOR (BBX-A7) (4
 K-PROTHORACICOTROPIC HORMONE) (4K-PTTH).//0.99:45:
 46//BOMBYX MORI (SILK MOTH).//P26730
 F-MAMMA1002937//ZINC FINGER PROTEIN 42 (MYELOID ZI
 NC FINGER 1) (MZF-1).//6.5e-24:147:34//HOMO SAPIEN
 S (HUMAN).//P28698

2957

F-MAMMA1002938//CERULOPLASMIN PRECURSOR (EC 1.16.3.1) (FERROXIDASE).//4.7e-11:44:68//MUS MUSCULUS (MOUSE).//Q61147

F-MAMMA1002941//PROTEIN Q300.//0.0076:21:61//MUS MUSCULUS (MOUSE).//Q02722

F-MAMMA1002947//HYPOTHETICAL 35.5 KD PROTEIN IN TRANSPOSON TN4556.//1.9e-08:152:38//STREPTOMYCES FRATIAE.//P20186

F-MAMMA1002964

F-MAMMA1002970//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//0.0057:55:43//HOMO SAPIENS (HUMAN).//P39189

F-MAMMA1002972//BRAIN-SPECIFIC HOMEBOX/POU DOMAIN PROTEIN 3A (BRN-3A) (OCT-T1) (HOMEBOX/POU DOMAIN PROTEIN RDC-1).//0.84:53:41//HOMO SAPIENS (HUMAN).//Q01851

F-MAMMA1002973//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!//4.6e-11:54:68//HOMO SAPIENS (HUMAN).//P39192

F-MAMMA1002982

F-MAMMA1002987//HYPOTHETICAL 11.9 KD PROTEIN IN RP C8-MFA2 INTERGENIC REGION.//0.17:47:29//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53906

F-MAMMA1003003//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//8.6e-09:30:73//HOMO SAPIENS (HUMAN).//P39195

F-MAMMA1003004//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//0.0071:41:58//HOMO SAPIENS (HUMAN).//P39195

F-MAMMA1003007//SPERM PROTAMINE P1.//0.0076:51:37//TACHYGLOSSUS ACULEATUS ACULEATUS (AUSTRALIAN ECHIDNA).//P35311

F-MAMMA1003011//HISTONE MACRO-H2A.1.//1.8e-60:175:70//RATTUS NORVEGICUS(RAT).//Q02874

F-MAMMA1003013//ACTIN BINDING PROTEIN.//0.097:83:31//SACCHAROMYCES EXIGUUS (YEAST).//P38479

F-MAMMA1003015

F-MAMMA1003019//MYOTUBULARIN.//0.022:56:37//HOMO SAPIENS (HUMAN).//Q13496

F-MAMMA1003026//HYPOTHETICAL 29.3 KD PROTEIN (ORF92).//0.0014:208:27//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPMPV).//O10341

F-MAMMA1003031//PROBABLE E4 PROTEIN (E1^E4).//0.14:49:32//HUMAN PAPILLOMAVIRUS TYPE 6B.//P06459

F-MAMMA1003035//HYPOTHETICAL 24.4 KD PROTEIN IN LP D 3' REGION (ORF4).//5.1e-12:112:34//ZYMOMONAS MOBILIS.//O66114

F-MAMMA1003039//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//2.4e-07:68:54//HOMO SAPIENS (HUMAN).//P39188

2958

F-MAMMA1003040//!!!! ALU SUBFAMILY SB1 WARNING ENTRY !!!!!//2.8e-39:90:57//HOMO SAPIENS (HUMAN).//P39190

F-MAMMA1003044

F-MAMMA1003047//SPERM HISTONE P2 PRECURSOR (PROTAMINE 2).//0.18:25:44//BOS TAURUS (BOVINE).//P19782

F-MAMMA1003049//PROBABLE E4 PROTEIN.//0.50:67:29//HUMAN PAPILLOMAVIRUS TYPE 6C.//P20969

F-MAMMA1003055//WEAK TOXIN CM-2.//0.99:23:30//NAJAH HAJE HAJE (EGYPTIAN COBRA).//P01415

F-MAMMA1003056//EXPORTED PROTEIN 7 (FRAGMENT).//1.0:52:32//STREPTOCOCCUS PNEUMONIAE.//P35597

F-MAMMA1003057//MD6 PROTEIN.//1.5e-85:168:95//MUS MUSCULUS (MOUSE).//Q60584

F-MAMMA1003066//REGB PROTEIN.//1.0:62:27//PSEUDOMONAS AERUGINOSA.//Q03381

F-MAMMA1003089//!!!! ALU SUBFAMILY SB1 WARNING ENTRY !!!!!//5.1e-15:44:77//HOMO SAPIENS (HUMAN).//P39190

20 F-MAMMA1003099//ENDOTHELIAL ACTIN-BINDING PROTEIN (ABP-280) (NONMUSCLE FILAMIN) (FILAMIN 1).//4.8e-20:80:62//HOMO SAPIENS (HUMAN).//P21333

F-MAMMA1003104//PHOTOSYSTEM I REACTION CENTRE SUBUNIT VIII.//0.98:22:40//SYNECHOCOCCUS ELONGATUS NAGELI.//P25900

F-MAMMA1003113//PROCOLLAGEN ALPHA 2(I) CHAIN PRECURSOR (FRAGMENTS).//0.67:35:45//GALLUS GALLUS (CHICKEN).//P02467

F-MAMMA1003127//MYOSIN I ALPHA (MMI-ALPHA).//5.2e-34:141:56//MUS MUSCULUS (MOUSE).//P46735

F-MAMMA1003135//HYPOTHETICAL 182.0 KD PROTEIN IN NMD5-HOM6 INTERGENIC REGION.//3.6e-05:91:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P47170

F-MAMMA1003140

F-MAMMA1003146//MALE SPECIFIC SPERM PROTEIN MST87 F.//1.0:33:36//DROSOPHILA MELANOGASTER (FRUIT FLY).//P08175

F-MAMMA1003150//HYPOTHETICAL 84.3 KD PROTEIN ZK945.10 IN CHROMOSOME II.//4.4e-10:254:30//CAENORHABDITIS ELEGANS.//Q09625

F-MAMMA1003166//BRAIN PROTEIN H5.//4.0e-42:182:48//HOMO SAPIENS (HUMAN).//O43236

F-NT2RM1000001//HYPOTHETICAL 8.7 KD PROTEIN IN RPL22-RPL23 INTERGENIC REGION (ORF70).//0.15:38:34//ASTASIA LONGA (EUGLENOPHYCEAN ALGA).//P34779

F-NT2RM1000018

F-NT2RM1000032//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (AGL).//0.51:17:41//CYPRINUS CARPIO (COMMON CARP).//P24948

50 F-NT2RM1000035//3-HYDROXY-3-METHYLGLUTARYL-COENZYME

2959

E A REDUCTASE (EC 1.1.1.34) (HMG-COA REDUCTASE). // 0.00011:114:27//BLATTELLA GERMANICA (GERMANCOCKROACH). //P54960
 F-NT2RM1000037//METALLOTHIONEIN-II (MT-II). //0.025:19:47//SCYLLA SERRATA(MUD CRAB). //P02806
 F-NT2RM1000039//VITELLINE MEMBRANE VM34CA PROTEIN PRECURSOR. //0.00083:84:33//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q06521
 F-NT2RM1000055//HISTIDINE-RICH GLYCOPROTEIN PRECURSOR. //1.1e-07:34:55//PLASMODIUM LOPHURAE. //P04929
 F-NT2RM1000059//MYOCYTE-SPECIFIC ENHANCER FACTOR 2 B (SERUM RESPONSE FACTOR-LIKE PROTEIN 2) (XMEF2) (RSRFR2). //0.18:83:36//HOMO SAPIENS (HUMAN). //Q02080
 F-NT2RM1000062//PROLINE-RICH PEPTIDE P-B. //0.54:34:44//HOMO SAPIENS (HUMAN). //P02814
 F-NT2RM1000080//HYPOTHETICAL 35.7 KD PROTEIN SLR1128. //2.1e-20:119:40//SYNECHOCYSTIS SP. (STRAIN PCC 6803). //P72655
 F-NT2RM1000086//SALIVARY PROLINE-RICH PROTEIN PO (ALLELE K) [CONTAINS: PEPTIDE P-D] (FRAGMENT). //0.20:56:35//HOMO SAPIENS (HUMAN). //P10162
 F-NT2RM1000092//COLLAGEN-LIKE PROTEIN. //0.0017:44:45//HERPESVIRUS SAIMIRI (SUBGROUP C / STRAIN 488). //P22576
 F-NT2RM1000118//CALCINEURIN B SUBUNIT (PROTEIN PHOSPHATASE 2B REGULATORY SUBUNIT) (CALCINEURIN REGULATORY SUBUNIT). //5.7e-07:109:28//NEUROSPORACRASSA. //P87072
 F-NT2RM1000119//TRANSCRIPTIONAL REGULATOR IE63 (VM W63) (ICP27). //0.0050:135:32//HERPES SIMPLEX VIRUS (TYPE 2 / STRAIN HG52). //P28276
 F-NT2RM1000127//EXTENSIN PRECURSOR (PROLINE-RICH GLYCOPROTEIN). //0.032:68:32//SORGHUM VULGARE (SORGHUM). //P24152
 F-NT2RM1000131//METALLOTHIONEIN-III (MT-III) (GROWTH INHIBITORY FACTOR) (GIF). //0.82:33:39//BOS TAURUS (BOVINE). //P37359
 F-NT2RM1000132//NADH-UBIQUINONE OXIDOREDUCTASE 13 KD-A SUBUNIT PRECURSOR (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-13KD-A) (CI-13KD-A). //2.7e-59:124:91//HOMO SAPIENS (HUMAN). //Q075380
 F-NT2RM1000153//CYTOSOLIC PURINE 5'-NUCLEOTIDASE (EC 3.1.3.5). //2.5e-08:148:29//HOMO SAPIENS (HUMAN). //P49902
 F-NT2RM1000186//CALCINEURIN B SUBUNIT (PROTEIN PHOSPHATASE 2B REGULATORY SUBUNIT) (CALCINEURIN REGULATORY SUBUNIT). //1.9e-07:109:27//NEUROSPORACRASSA. //P87072
 F-NT2RM1000187//PUTATIVE PRE-MRNA SPLICING FACTOR

2960

ATP-DEPENDENT RNA HELICASE SPAC10F6.02C. //1.0e-12:94:46//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //042643
 F-NT2RM1000199//CUTICLE COLLAGEN 12 PRECURSOR. //0.46:130:33//CAENORHABDITIS ELEGANS. //P20630
 F-NT2RM1000242//PUTATIVE ATP SYNTHASE J CHAIN, MITOCHONDRIAL (EC 3.6.1.34). //0.85:38:36//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q13931
 F-NT2RM1000244//HYPOTHETICAL 131.5 KD PROTEIN C02F12.7 IN CHROMOSOME X. //0.0055:98:36//CAENORHABDITIS ELEGANS. //Q11102
 F-NT2RM1000252//TRICHOHYALIN. //2.9e-06:88:36//OVIS ARIES (SHEEP). //P22793
 F-NT2RM1000256//GLUCOSAMINE--FRUCTOSE-6-PHOSPHATE AMINOTRANSFERASE [ISOMERIZING] (EC 2.6.1.16) (HEXOSEPHOSPHATE AMINOTRANSFERASE) (D-FRUCTOSE-6-PHOSPHATE AMINOTRANSFERASE) (GFAT). //2.9e-54:153:67//MUS MUSCULUS (HOUSE). //P47856
 F-NT2RM1000257//MAGO NASHI PROTEIN. //5.9e-64:136:89//DROSOPHILA MELANOGASTER (FRUIT FLY). //P49028
 F-NT2RM1000260
 F-NT2RM1000271//GALACTOKINASE (EC 2.7.1.6). //0.99:41:39//BACILLUS SUBTILIS. //P39574
 F-NT2RM1000272//HYPOTHETICAL 55.5 KD PROTEIN ZK1128.2 IN CHROMOSOME III. //8.8e-25:131:45//CAENORHABDITIS ELEGANS. //Q09357
 F-NT2RM1000280//VACUOLAR ATP SYNTHASE SUBUNIT D (EC 3.6.1.34) (V-ATPASE SUBUNIT) (V-ATPASE 28 KD ACCESSORY PROTEIN). //2.5e-63:121:94//BOS TAURUS (BOVINE). //P39942
 F-NT2RM1000300//TREACLE PROTEIN (TREACHER COLLINS SYNDROME PROTEIN). //0.51:145:26//HOMO SAPIENS (HUMAN). //Q13428
 F-NT2RM1000314
 F-NT2RM1000318//50S RIBOSOMAL PROTEIN L23. //0.83:28:35//AQUIFEX AEOLICUS. //Q066433
 F-NT2RM1000341
 F-NT2RM1000354//HYPOTHETICAL 5.8 KD PROTEIN IN PUBA 5' REGION (ORF55). //0.95:43:37//RHODOBACTER CAPSULATUS (RHODOPSEUDOMONAS CAPSULATA). //P26159F-NT2RM1000355//SPERM-SPECIFIC PROTEIN PHI-1. //0.0016:73:43//MYTILUS EDULIS (BLUE MUSSEL). //Q04621
 F-NT2RM1000365//HYPOTHETICAL PROTEIN KIAA0140. //3.5e-10:83:49//HOMO SAPIENS (HUMAN). //Q14153
 F-NT2RM1000377//DUAL SPECIFICITY PROTEIN PHOSPHATASE 9 (EC 3.1.3.48) (EC3.1.3.16) (MITOGEN-ACTIVATED PROTEIN KINASE PHOSPHATASE 4) (MAP KINASE PHOSPHATASE 4) (MKP-4). //4.9e-18:113:38//HOMO SAPIENS (HUMAN). //Q99956
 F-NT2RM1000388//HYPOTHETICAL 27.7 KD PROTEIN IN CP

T1-SPC98 INTERGENIC REGION.//0.00023:67:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53915
 F-NT2RM1000394//HISTONE H3.3 (H3.B) (H3.3Q).//4.7e-52:71:91//HOMO SAPIENS (HUMAN), MUS MUSCULUS (MOUSE), RATTUS NORVEGICUS (RAT), ORYCTOLAGUS CUNICULUS (RABBIT), GALLUS GALLUS (CHICKEN), SPISULA SOLIDISSIMA (ATLANTIC SURF-CLAM), DROSOPHILA MELANOGASTER (FRUIT FLY), AND DROSOPHILA HYDEI (FRUIT FLY).//P06351
 F-NT2RM1000399//ENDOTHELIN-2 PRECURSOR (ET-2) (FRAGMENT).//0.92:24:45//CANIS FAMILIARIS (DOG).//P12064
 F-NT2RM1000421//CUTICLE COLLAGEN 2C (FRAGMENT).//0.12:93:33//HAEMONCHUS CONTORTUS.//P16252
 F-NT2RM1000430//PISTIL-SPECIFIC EXTENSIN-LIKE PROTEIN PRECURSOR (PELP).//0.13:86:31//NICOTIANA TABACUM (COMMON TOBACCO).//Q03211
 F-NT2RM1000499//HYPOTHETICAL PROTEIN KIAA0041 (FRAGMENT).//2.9e-17:75:49//HOMO SAPIENS (HUMAN).//Q15057
 F-NT2RM1000539//HYPOTHETICAL 10.4 KD PROTEIN IN FT R1-SPT15 INTERGENIC REGION.//2.9e-16:82:51//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40089
 F-NT2RM1000553//GLYCOLIPID TRANSFER PROTEIN (GLTP).//6.4e-06:103:33//SUSSCROFA (PIG).//P17403
 F-NT2RM1000555//UNR PROTEIN.//8.7e-77:105:95//RATTUS NORVEGICUS (RAT).//P18395
 F-NT2RM1000563//DNA-BINDING P52/P100 COMPLEX, 100 KD SUBUNIT (FRAGMENTS).//0.15:20:50//HOMO SAPIENS (HUMAN).//P30808
 F-NT2RM1000623//CLARA CELL PHOSPHOLIPID-BINDING PROTEIN PRECURSOR (CCBPB) (CLARA CELLS 10 KD SECRETORY PROTEIN) (CC10).//0.17:70:34//HOMO SAPIENS (HUMAN).//P11684
 F-NT2RM1000648//GLYCOSYLTRANSFERASE ALG2 (EC 2.4.1.-).//2.0e-22:133:42//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P43636
 F-NT2RM1000661//METALLOTHIONEIN-III (MT-III) (GROWTH INHIBITORY FACTOR) (GIF) (GIFB).//0.0060:24:33//HOMO SAPIENS (HUMAN).//P25713
 F-NT2RM1000666//COLD SHOCK PROTEIN SCOF.//9.1e-07:67:41//STREPTOMYCES COELICOLOR.//P48859
 F-NT2RM1000669//CHLOROPLAST 50S RIBOSOMAL PROTEIN L31.//0.071:69:31//PORPHYRA PURPUREA.//P51290
 F-NT2RM1000672//SIGNAL RECOGNITION PARTICLE SEC65 SUBUNIT (FRAGMENT).//0.27:42:42//KLUYVEROMYCES LACTIS (YEAST).//013475
 F-NT2RM1000691//RETINOBLASTOMA BINDING PROTEIN 2 (RBBP-2).//4.3e-42:241:42//HOMO SAPIENS (HUMAN).//P29375

F-NT2RM1000699//N2,N2-DIMETHYLGUANOSINE TRNA METHYLTRANSFERASE PRECURSOR (EC 2.1.1.32).//0.94:48:37//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P15565
 F-NT2RM1000702//GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT 1.//0.0013:139:25//DROSOPHILA MELANOGASTER (FRUIT FLY).//P26308
 F-NT2RM1000725//BASIC PROLINE-RICH PEPTIDE P-E (IB-9).//1.0:15:60//HOMO SAPIENS (HUMAN).//P02811
 F-NT2RM1000741//STATHMIN (CLONE X020) (FRAGMENT).//1.0:53:32//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//Q09005
 F-NT2RM1000742//HYPOTHETICAL 24.1 KD PROTEIN IN DHFR 3' REGION (ORF2).//1.0:54:42//HERPESVIRUS SAIMIRI (STRAIN 484-77).//P25049
 F-NT2RM1000746//HYPOTHETICAL 16.8 KD PROTEIN C29E 6.04 IN CHROMOSOME I.//0.11:87:21//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09858
 F-NT2RM1000770//DXS6673E PROTEIN.//2.0e-38:190:48//HOMO SAPIENS (HUMAN).//Q14202
 20 F-NT2RM1000772//VEGETATIBLE INCOMPATIBILITY PROTEIN HET-E-1.//4.3e-12:141:30//PODOSPORA ANSERINA.//Q00808
 F-NT2RM1000780//MALE SPECIFIC SPERM PROTEIN MST87 F.//0.98:34:38//DROSOPHILA MELANOGASTER (FRUIT FLY).//P08175
 F-NT2RM1000781
 F-NT2RM1000800//24.1 KD PROTEIN IN VMA12-APN1 INTERGENIC REGION.//7.9e-11:135:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P28707
 30 F-NT2RM1000802//ALPHA-AMYLASE INHIBITOR PAM I (PIG PANCREATIC ALPHA-AMYLASE INHIBITOR OF MICROBES I).//0.43:62:35//STREPTOMYCES OLIVACEOVIRIDIS (STREPTOMYCES CORCHORUSII).//P09921
 F-NT2RM1000811
 F-NT2RM1000826//UNR PROTEIN.//1.1e-110:144:83//RATTUS NORVEGICUS (RAT).//P18395
 F-NT2RM1000829//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (AGL).//1.0:38:34//DROSOPHILA SIMULANS (FRUIT FLY).//P50270
 40 F-NT2RM1000833//PROTEIN TRANSPORT PROTEIN SEC61 ALPHA SUBUNIT.//1.4e-62:145:84//CANIS FAMILIARIS (DOG).//P38377
 F-NT2RM1000850//TESTIS-SPECIFIC PROTEIN KINASE 1 (EC 2.7.1.-).//6.1e-08:136:33//RATTUS NORVEGICUS (RAT).//Q63572
 F-NT2RM1000852//ATP-DEPENDENT RNA HELICASE ROK1.//1.6e-34:212:43//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P45818
 F-NT2RM1000857//HISTONE H1.6.1.//0.76:31:48//TRYPAANOSOMA CRUZI.//P40273

2963

F-NT2RM1000867//MICROSOMAL SIGNAL PEPTIDASE 10.8 K D SUBUNIT (EC 3.4.-.-).//0.0082:76:25//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P46965
 F-NT2RM1000874//DNA-BINDING P52/P100 COMPLEX, 100 KD SUBUNIT (FRAGMENTS).//0.38:12:58//HOMO SAPIENS (HUMAN).//P30808
 F-NT2RM1000882//CYTOCHROME B5.//9.0e-13:92:38//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40312
 F-NT2RM1000883//DNA-BINDING P52/P100 COMPLEX, 100 KD SUBUNIT (FRAGMENTS).//0.79:22:59//HOMO SAPIENS (HUMAN).//P30808
 F-NT2RM1000885//HYPOTHETICAL 5.8 KD PROTEIN.//0.76:18:38//CLOVER YELLOW MOSAIC VIRUS (CYMV).//P16485
 F-NT2RM1000894//DNA-DIRECTED RNA POLYMERASE I 135 KD POLYPEPTIDE (EC 2.7.7.6) (RNA POLYMERASE I SUBUNIT 2) (RPA135) (RNA POLYMERASE I 127 KD SUBUNIT).//6.2e-70:153:88//RATTUS NORVEGICUS (RAT).//054888
 F-NT2RM1000898//ACTIN, CYTOPLASMIC (ACTIN, MICRONUCLEAR).//4.3e-12:159:28//OXYTRICHA FALLAX.//P02583
 F-NT2RM1000905//GLUTATHIONE S-TRANSFERASE 1-1 (EC 2.5.1.18) (CLASS-THETA).//0.98:39:35//LUCILIA CUPRINA (GREENBOTTLE FLY) (AUSTRALIAN SHEEP BLOWFLY).//P42860
 F-NT2RM1000924//HYPOTHETICAL 39.7 KD PROTEIN C34E1.0.2 IN CHROMOSOME III.//1.3e-11:169:28//CAENORHABDITIS ELEGANS.//P46577
 F-NT2RM1000927//CUTICLE COLLAGEN 1.//0.00048:141:31//CAENORHABDITIS ELEGANS.//P08124
 F-NT2RM1000962//HYPOTHETICAL 35.8 KD PROTEIN C4F8.04 IN CHROMOSOME I.//7.1e-13:169:31//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//014180
 F-NT2RM1000978//HYPOTHETICAL 20.2 KD PROTEIN IN MN4-PTK1 INTERGENIC REGION.//0.61:82:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P36045
 F-NT2RM1001003//ALPHA-2 CATENIN (ALPHA N-CATENIN) (NEURAL ALPHA-CATENIN).//1.6e-21:211:31//GALLUS GALLUS (CHICKEN).//P30997
 F-NT2RM1001008//HYPOTHETICAL 72.5 KD PROTEIN C2F7.10 IN CHROMOSOME I.//3.2e-15:119:36//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09701
 F-NT2RM1001043//ENDOTHELIN-1 (ET-1) (FRAGMENT).//0.78:32:34//MACACA FASCICULARIS (CRAB EATING MACAQUE) (CYNOMOLGUS MONKEY).//Q28469
 F-NT2RM1001044
 F-NT2RM1001059//LORICRIN.//8.6e-08:108:39//HOMO SAPIENS (HUMAN).//P23490
 F-NT2RM1001066//METALLOTHIONEIN-LIKE PROTEIN TYPE 2.//0.99:24:50//LYCOPERSICON ESCULENTUM (TOMATO).//Q43513
 F-NT2RM1001072//1-PHOSPHATIDYLINOSITOL-4,5-BISPHOS

2964

PHATE PHOSPHODIESTERASE GAMMA 1 (EC 3.1.4.11) (PLC-GAMMA-1) (PHOSPHOLIPASE C-GAMMA-1) (PLC-II) (PLC-148).//4.7e-15:148:33//HOMO SAPIENS (HUMAN).//P19174
 F-NT2RM1001074//HYPOTHETICAL PROTEIN F-215.//8.6e-05:126:30//HUMAN ADENOVIRUS TYPE 2.//P03291
 F-NT2RM1001082//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//6.5e-19:75:54//HOMO SAPIENS (HUMAN).//P39195
 F-NT2RM1001085//MALE SPECIFIC SPERM PROTEIN MST84DB.//0.49:29:41//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01643
 F-NT2RM1001092//ZINC FINGER PROTEIN 83 (ZINC FINGER PROTEIN HPF1).//2.8e-42:200:38//HOMO SAPIENS (HUMAN).//P51522
 F-NT2RM1001102//HYPOTHETICAL 62.8 KD PROTEIN IN TA F145-YOR1 INTERGENIC REGION.//1.7e-18:161:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53331
 F-NT2RM1001105//HYPOTHETICAL 35.5 KD PROTEIN IN TRANSPOSON TN4556.//4.0e-05:157:35//STREPTOMYCES FRADIACE.//P20186
 F-NT2RM1001112//NONHISTONE CHROMOSOMAL PROTEIN HMG-17.//0.18:20:55//BOSTAURUS (BOVINE).//P02313
 F-NT2RM1001115
 F-NT2RM1001139//GLYCINE-RICH CELL WALL STRUCTURAL PROTEIN 1.8 PRECURSOR (GRP 1.8).//2.0e-25:156:46//PASEOLUS VULGARIS (KIDNEY BEAN) (FRENCH BEAN).//P10496
 F-NT2RM2000006//MITOCHONDRIAL RIBOSOMAL PROTEIN S12.//0.76:45:35//LEISHMANIA TARENTOLAE (SAUROLEISHMANIA TARENTOLAE).//Q34940
 F-NT2RM2000013//DNA-DIRECTED RNA POLYMERASE III 128 KD POLYPEPTIDE (EC 2.7.7.6) (RNA POLYMERASE III SUBUNIT 2).//3.9e-87:238:65//DROSOPHILA MELANOGASTER (FRUIT FLY).//P25167
 F-NT2RM2000030//TOXINS 1 AND 2.//0.98:21:42//TRIMERESURUS WAGLERI (WAGLER'S PIT VIPER) (TROPIDOLAEMUS WAGLERI).//P24335
 F-NT2RM2000032//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//0.00059:53:49//HOMO SAPIENS (HUMAN).//P39188
 F-NT2RM2000042//SMALL PROLINE-RICH PROTEIN II (SPR-II) (CLONE 174N).//1.0:68:26//HOMO SAPIENS (HUMAN).//P22532
 F-NT2RM2000092//HYPOTHETICAL 67.5 KD PROTEIN IN PRPS4-STE20 INTERGENIC REGION.//7.0e-11:80:40//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38748F-NT2RM2000093//OVARY MATURATING PARSIN (OMP).//1.0:26:38//LOCUSTA MIGRATORIA (MIGRATORY LOCUST).//P80045
 F-NT2RM2000101//HYPOTHETICAL 39.3 KD PROTEIN C02B

2966

- 8.6 IN CHROMOSOME X. //3.3e-09: 56: 35//CAENORHABDITI
S ELEGANS. //Q11096
F-NT2RM2000124//CAMP-DEPENDENT PROTEIN KINASE, ALP
HA-CATALYTIC SUBUNIT (EC 2.7.1.37) (PKA C-ALPHA). /
/3.1e-35: 77: 96//MUS MUSCULUS (MOUSE). //P05132
F-NT2RM2000191//3', 5'-CYCLIC-NUCLEOTIDE PHOSPHODIE
STERASE REGA (EC 3.1.4.17) (PDEASE REGA). //3.3e-0
5: 181: 27//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD). //
Q23917
F-NT2RM2000192//REPLICATION PROTEIN E1 (FRAGMENT 10
S). //0.019: 148: 25//COTTONTAIL RABBIT (SHOPE) PAPIL
LOMAVIRUS (STRAIN WASHINGTON B) (CRPV). //P51894
F-NT2RM2000239//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T). //0.00032: 111: 32//MUS MUSCULUS (MOUSE). //P05143
F-NNNNNNNNNNNN//METALLOTHIONEIN-LIKE PROTEIN TYPE
2. //0.046: 59: 33//LYCOPERSICON ESCULENTUM (TOMATO).
//Q43512
F-NT2RM2000250//GALECTIN-3 (GALACTOSE-SPECIFIC LEC
TIN 3) (MAC-2 ANTIGEN) (IGE-BINDING PROTEIN) (35 KD
LECTIN) (CARBOHYDRATE BINDING PROTEIN 35) (CBP 35) 20
(LAMININ-BINDING PROTEIN) (LECTIN L-29). //0.054: 4
6: 34//RATTUS NORVEGICUS (RAT). //P08699
F-NT2RM2000259//TRANS-ACTING TRANSCRIPTIONAL PROTE
IN ICPO (P135 PROTEIN) (IER 2.9/ER2.6). //0.27: 112: 3
3//BOVINE HERPESVIRUS TYPE 1 (STRAIN JURA). //P2912
8
F-NT2RM2000260//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T). //4.7e-22: 191: 35//MUS MUSCULUS (MOUSE). //P05143
F-NT2RM2000287//HYPOTHETICAL 11.8 KD PROTEIN C1B3.
02C IN CHROMOSOME I. //5.0e-19: 83: 53//SCHIZOSACCHAR 30
OMYCES POMBE (FISSION YEAST). //013868
F-NT2RM2000322//DIAMINOPELMATE DECARBOXYLASE (EC
4.1.1.20) (DAP DECARBOXYLASE). //0.47: 117: 29//HELIC
OBACTER PYLORI (CAMPYLOBACTER PYLORI). //P56129
F-NT2RM2000359//SPORE GERMINATION PROTEIN 270-11. /
/0.12: 83: 36//DICTYOSTELIUM DISCOIDEUM (SLIME MOL
D). //P22698
F-NT2RM2000363//BREAKPOINT CLUSTER REGION PROTEIN.
//1.3e-16: 203: 30//HOMOSAPIENS (HUMAN). //P11274
F-NT2RM2000368//DEK PROTEIN. //0.00027: 100: 32//HOMO 40
SAPIENS (HUMAN). //P35659
F-NT2RM2000371//POLYRIBONUCLEOTIDE NUCLEOTIDYLTRAN
SFERASE (EC 2.7.7.8) (POLYNUCLEOTIDE PHOSPHORYLAS
E) (PNPASE). //6.8e-36: 170: 47//ESCHERICHIA COLI. //P
05055
F-NT2RM2000374//NODAL PRECURSOR. //1.1e-32: 64: 95//M
US MUSCULUS (MOUSE). //P43021
F-NT2RM2000395//IMMEDIATE-EARLY PROTEIN IE180. //0.
31: 41: 43//PSEUDORABIESVIRUS (STRAIN INDIANA-FUNKHA
USER / BECKER) (PRV). //P11675
F-NT2RM2000402//ENDOSOMAL P24A PROTEIN PRECURSOR
(70 KD ENOMEMBRANE PROTEIN) (PHEROMONE ALPHA-FACT
OR TRANSPORTER) (ACIDIC 24 KD LATE ENDOCYTIICINTERM
EDIATE COMPONENT). //1.2e-30: 228: 32//SACCHAROMYCES
CEREVISIAE (BAKER'S YEAST). //P32802
F-NT2RM2000407//TRANSMEMBRANE PROTEIN SEX PRECURSO
R. //0.032: 105: 30//HOMOSAPIENS (HUMAN). //P51805
F-NT2RM2000420//METALLOTHIONEIN (MT). //0.88: 42: 38/
/PLEURONECTES PLATESSA (PLAICE). //P07216
F-NT2RM2000422//SODIUM- AND CHLORIDE-DEPENDENT TRA
NSPORTER NTT73. //2.0e-117: 237: 87//RATTUS NORVEGICU
S (RAT). //Q08469
F-NT2RM2000452//HYPOTHETICAL 63.6 KD PROTEIN IN YP
T52-GCN3 INTERGENIC REGION. //1.1e-08: 157: 28//SACCH
AROMYCES CEREVISIAE (BAKER'S YEAST). //P36113F-NT2R
M2000469//70 KD ANTIGEN. //0.050: 207: 23//SHIGELLA F
LEXNERI. //P18010F-NT2RM2000490//BASIC PROLINE-RICH
PEPTIDE P-E (IB-9). //0.022: 25: 44//HOMO SAPIENS (H
UMAN). //P02811
F-NT2RM2000502//MALE SPECIFIC SPERM PROTEIN MST84D
D. //0.0037: 17: 58//DROSOPHILA MELANOGASTER (FRUIT F
LY). //Q01645
F-NT2RM2000504//HYPOTHETICAL 99.0 KD PROTEIN SPBC1
19.17. //1.7e-22: 195: 27//SCHIZOSACCHAROMYCES POMBE
(FISSION YEAST). //042908
F-NT2RM2000522//RAS-RELATED PROTEIN RABA (FRAGMEN
T). //3.6e-05: 67: 29//DICTYOSTELIUM DISCOIDEUM (SLIM
E MOLD). //P34141
F-NT2RM2000540//HYPOTHETICAL 83.8 KD PROTEIN C27F
2.7 IN CHROMOSOME III. //8.4e-33: 214: 38//CAENORHABD 30
ITIS ELEGANS. //Q18262
F-NT2RM2000556//HYPOTHETICAL PROTEIN KIAA0288 (HA6
116). //1.7e-09: 133: 36//HOMO SAPIENS (HUMAN). //P565
24
F-NT2RM2000566//INTEGRIN ALPHA-6 PRECURSOR (VLA-6)
(CD49F). //2.2e-60: 244: 51//HOMO SAPIENS (HUMAN). //P
23229
F-NT2RM2000567//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T). //2.3e-09: 192: 34//MUS MUSCULUS (MOUSE). //P05143
F-NT2RM2000569//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//9.0e-08: 43: 72//HOMO SAPIENS (HUMAN). //P3918
8
F-NT2RM2000577//ISOLEUCYL-TRNA SYNTHETASE (EC 6.1.
1.5) (ISOLEUCINE--TRNALIGASE) (ILERS). //9.1e-54: 22
5: 45//SYNECHOCYSTIS SP. (STRAIN PCC 6803). //P73505
F-NT2RM2000581//SPLICEOSOME ASSOCIATED PROTEIN 49
(SAP 49) (SF3B53). //0.079: 111: 34//HOMO SAPIENS (HU
MAN). //Q15427
F-NT2RM2000588//HYPOTHETICAL PROTEIN KIAA0288 (HA6
116). //2.3e-09: 193: 32//HOMO SAPIENS (HUMAN). //P565
50

24

F-NT2RM2000594//BASIC PROLINE-RICH PEPTIDE P-E (1B-9). //0.18:33:42//HOMOSAPIENS (HUMAN). //P02811

F-NT2RM2000599//DNA (CYTOSINE-5)-METHYLTRANSFERASE (EC 2.1.1.37) (DNA METHYLTRANSFERASE) (DNA METASE) (MCMT) (M. MMUI). //1.5e-09:68:45//MUS MUSCULUS (MOUSE). //P13864

F-NT2RM2000609//GRANULIN 2. //0.83:42:35//CYPRIUS CARPIO (COMMON CARP). //P81014

F-NT2RM2000612//ZINC FINGER PROTEIN GCS1. //7.2e-05:155:29//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P35197

F-NT2RM2000623//A-AGGLUTININ ATTACHMENT SUBUNIT PRECURSOR. //1.8e-09:196:33//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32323

F-NT2RM2000624//SALIVARY GLUE PROTEIN SGS-3 PRECURSOR. //0.070:113:27//DROSOPHILA ERECTA (FRUIT FLY). //P13730

F-NT2RM2000635//SPERM PROTAMINE P1. //0.54:47:38//A NTECHINUS STUARTII. //P42129

F-NT2RM2000636//OUTER MEMBRANE PROTEIN H.8 PRECURSOR. //0.096:62:35//NEISSERIA GONORRHOEA. //P11910

F-NT2RM2000639//HYPOTHETICAL PROTEIN MJ0243. //0.99:32:34//METHANOCOCCUS JANNASCHII. //Q57694

F-NT2RM2000649//NEURONAL CALCIUM SENSOR 1 (NCS-1). //0.00049:70:35//RATTUS NORVEGICUS (RAT), AND GALLUS GALLUS (CHICKEN). //P36610

F-NT2RM2000669//50S RIBOSOMAL PROTEIN L34. //1.0:34:44//BACILLUS SUBTILIS. //P05647

F-NT2RM2000691//ACTIN-LIKE PROTEIN 3 (ACTIN-2). //7.0e-116:243:87//HOMO SAPIENS (HUMAN), AND BOS TAURUS (BOVINE). //P32391

F-NT2RM2000714//RAS SUPPRESSOR PROTEIN 1 (RSU-1) (RSP-1 PROTEIN) (RSP-1). //3.8e-21:174:35//HOMO SAPIENS (HUMAN). //Q15404

F-NT2RM2000718//HYPOTHETICAL 52.9 KD SERINE-RICH PROTEIN C11C7.01 IN CHROMOSOME I. //0.0022:174:29//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //013695

F-NT2RM2000735//ZINC FINGER PROTEIN 43 (ZINC PROTEIN HTF6). //1.6e-102:246:74//HOMO SAPIENS (HUMAN). //P28160

F-NT2RM2000740//HYPOTHETICAL 131.1 KD HELICASE IN ALG7-ENP1 INTERGENIC REGION. //8.5e-51:212:49//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P38144

F-NT2RM2000795//ALU SUBFAMILY SB WARNING ENTRY !!!!! //9.0e-41:125:53//HOMO SAPIENS (HUMAN). //P39189

F-NT2RM2000821//COATOMER BETA SUBUNIT (BETA-COAT PROTEIN) (BETA-COP). //1.1e-128:291:89//RATTUS NORVEGICUS (RAT). //P23514

F-NT2RM2000837//CYCLIN-DEPENDENT KINASE INHIBITOR 1C (CYCLIN-DEPENDENT KINASE INHIBITOR P57) (P57KIP2). //3.9e-05:113:36//HOMO SAPIENS (HUMAN). //P49918

F-NT2RM2000951//HYPOTHETICAL 60.3 KD PROTEIN R08D 7.7 IN CHROMOSOME III. //2.5e-49:273:39//CAENORHABDITIS ELEGANS. //P30646

F-NT2RM2000952//NEUROFILAMENT TRIPLET H PROTEIN (200 KD NEUROFILAMENT PROTEIN) (NF-H) (FRAGMENT). //0.037:234:23//RATTUS NORVEGICUS (RAT). //P16884

10 F-NT2RM2000984//HYPOTHETICAL 54.7 KD PROTEIN F37A 4.1 IN CHROMOSOME III. //6.3e-44:216:43//CAENORHABDITIS ELEGANS. //P41879

F-NT2RM2001004//SYNAPSINS IA AND IB. //0.15:178:32//RATTUS NORVEGICUS (RAT). //P09951

F-NT2RM2001035//CCR4-ASSOCIATED FACTOR 1 (CAF1). //1.4e-87:188:90//MUS MUSCULUS (MOUSE). //Q60809

F-NT2RM2001065//ATP SYNTHASE A CHAIN (EC 3.6.1.34) (PROTEIN 6). //0.53:122:31//TRYPANOSOMA BRUCEI BRUCEI. //P24499

20 F-NT2RM2001100//HYPOTHETICAL 39.7 KD PROTEIN C34E1 0.2 IN CHROMOSOME III. //3.4e-13:171:30//CAENORHABDITIS ELEGANS. //P46577

F-NT2RM2001105//SPORE COAT PROTEIN SP96. //7.8e-06:141:34//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD). //P14328

F-NT2RM2001131//PROBABLE EUKARYOTIC INITIATION FACTOR C17C9.03. //2.3e-18:249:31//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10475

30 F-NT2RM2001141//HYPOTHETICAL 115.4 KD PROTEIN ZK75 7.3 IN CHROMOSOME III. //0.050:134:26//CAENORHABDITIS ELEGANS. //P34681

F-NT2RM2001152

F-NT2RM2001177//COLLAGEN ALPHA 1(XIV) CHAIN PRECURSOR (UNDULIN). //0.86:42:40//GALLUS GALLUS (CHICKEN). //P32018

F-NT2RM2001194//SMOOTH ELIN. //4.7e-05:77:32//HOMO SAPIENS (HUMAN). //P53814

F-NT2RM2001196//PROLINE-RICH PROTEIN MP-3 (FRAGMENT). //1.7e-18:218:35//MUS MUSCULUS (MOUSE). //P05143

40 F-NT2RM2001201//CYSTEINE STRING PROTEIN (CCCS1). //0.041:22:59//TORPEDO CALIFORNICA (PACIFIC ELECTRIC RAY). //P56101

F-NT2RM2001221//KALIRIN (PAM COOH-TERMINAL INTERACTOR PROTEIN 10) (P-CIP10). //1.3e-13:183:32//RATTUS NORVEGICUS (RAT). //P97924

F-NT2RM2001238//GLUTAMINASE, KIDNEY ISOFORM PRECURSOR (EC 3.5.1.2) (GLS) (L-GLUTAMINE AMIDOHYDROLASE). //6.5e-121:218:98//RATTUS NORVEGICUS (RAT). //P13264

50 F-NT2RM2001243//HYPOTHETICAL 200.0 KD PROTEIN IN G

2969

ZF3-IME2 INTERGENIC REGION. //0.00019:177:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P42945
 F-NT2RM2001247//LEGUMIN B (FRAGMENT). //0.22:54:35//PISUM SATIVUM (GARDENPEA). //P14594
 F-NT2RM2001256//PROTEIN TSG24 (MEIOTIC CHECK POINT REGULATOR). //1.8e-109:207:98//MUS MUSCULUS (MOUSE). //P53995
 F-NT2RM2001291//SMALL PROLINE RICH PROTEIN II (SPR-11) (CLONE 930). //0.016:22:40//HOMO SAPIENS (HUMAN). //P22531
 F-NT2RM2001306//REF(2)P PROTEIN. //0.61:51:33//DROSOPHILA MELANOGASTER (FRUIT FLY). //P14199
 F-NT2RM2001312//ALU SUBFAMILY SX WARNING ENTRY 11111//7.2e-11:33:72//HOMO SAPIENS (HUMAN). //P39195
 F-NT2RM2001319
 F-NT2RM2001324//ZYXIN. //5.1e-22:91:38//GALLUS GALLUS (CHICKEN). //Q04584
 F-NT2RM2001345//VEGETATILE INCOMPATIBILITY PROTEIN HET-E-1. //7.4e-10:159:27//PODOSPORA ANSERINA. //Q00808
 F-NT2RM2001360//ACCESSORY GLAND PEPTIDE PRECURSOR (PARAGONIAL PEPTIDE B). //1.0:27:48//DROSOPHILA MELANOGASTER (FRUIT FLY). //P05623
 F-NT2RM2001370//NAPE PROTEIN. //0.98:44:31//PARACOCUS DENITRIFICANS (SUBSP. THIOSPHAERA PANTOTROPH A). //Q56348
 F-NT2RM2001393//VITELLOGENIN PRECURSOR (VTC) [CONTAINS: LIPOVITELLIN LV-1N; LIPOVITELLIN LV-1C; LIPOVITELLIN LV-2]. //0.0024:163:31//ICHTHYOMYZONUNICUS PUS (SILVER LAMPREY). //Q91062
 F-NT2RM2001420
 F-NT2RM2001424//HETEROGENOUS NUCLEAR RIBONUCLEOPROTEIN U (HNRNP U). //2.4e-41:140:59//HOMO SAPIENS (HUMAN). //Q00839
 F-NT2RM2001499//HIGH-AFFINITY CATIONIC AMINO ACID TRANSPORTER-1 (CAT-1) (CAT1) (SYSTEM Y+ BASIC AMINO ACID TRANSPORTER) (ECOTROPIC RETROVIRAL LEUKEMIA RECEPTOR HOMOLOG) (ERR) (ECOTROPIC RETROVIRUS RECEPTOR HOMOLOG). //3.7e-71:201:68//HOMO SAPIENS (HUMAN). //P30825
 F-NT2RM2001504//CUTICLE COLLAGEN 2. //0.028:41:39//CAENORHABDITIS ELEGANS. //P17656
 F-NT2RM2001524//HYPOTHETICAL 61.3 KD PROTEIN F25B 5.5 IN CHROMOSOME III. //6.7e-47:190:42//CAENORHABDITIS ELEGANS. //Q09316
 F-NT2RM2001544//TELOMERE-BINDING PROTEIN 51 KD SUBUNIT. //0.0027:136:33//EUPLOTES CRASSUS. //Q06184
 F-NT2RM2001547//HYPOTHETICAL 48.6 KD PROTEIN IN BE T1-PAN1 INTERGENIC REGION. //8.5e-18:91:50//SACCHAR

2970

OMYCES CEREVISIAE (BAKER'S YEAST). //P40564
 F-NT2RM2001575//52 KD RO PROTEIN (SJOGREN SYNDROME TYPE A ANTIGEN (SS-A)) (RO(SS-A)). //3.9e-35:212:41//HOMO SAPIENS (HUMAN). //P19474
 F-NT2RM2001582//RESA PROTEIN. //0.0033:72:27//BACILLUS SUBTILIS. //P35160
 F-NT2RM2001588//EXTENSIN PRECURSOR (PROLINE-RICH GLYCOPROTEIN). //1.0e-06:115:32//ZEA MAYS (MAIZE). //P14918
 10 F-NT2RM2001592//KERATIN, ULTRA HIGH-SULFUR MATRIX PROTEIN (UHS KERATIN). //0.033:156:23//HOMO SAPIENS (HUMAN). //P26371
 F-NT2RM2001605//RETINOBLASTOMA BINDING PROTEIN 2 (RBBP-2). //1.1e-116:249:82//HOMO SAPIENS (HUMAN). //P29375
 F-NT2RM2001613//PROTEIN TRANSPORT PROTEIN SEC61 ALPHA SUBUNIT. //1.2e-97:192:100//RATTUS NORVEGICUS (RAT). //P38378
 F-NT2RM2001632//A-AGGLUTININ ATTACHMENT SUBUNIT PRECURSOR. //0.00068:145:28//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32323
 20 F-NT2RM2001635//NUCLEAR ENVELOPE PORE MEMBRANE PROTEIN POM 121 (PORE MEMBRANE PROTEIN OF 121 KD) (P145). //1.1e-39:235:47//RATTUS NORVEGICUS (RAT). //P52591
 F-NT2RM2001637//HYPOTHETICAL BHLF1 PROTEIN. //0.075:197:29//EPSTEIN-BARRVIRUS (STRAIN B95-8) (HUMAN HERPESVIRUS 4). //P03181
 F-NT2RM2001641//NADH-CYTOCHROME B5 REDUCTASE (EC 1.6.2.2) (B5R). //0.013:29:68//HOMO SAPIENS (HUMAN). //P00387
 F-NT2RM2001648//PROTEIN TRANSPORT PROTEIN SEC61 ALPHA SUBUNIT. //3.2e-65:132:100//CANIS FAMILIARIS (DOG). //P38377
 F-NT2RM2001652//PROTEIN TRANSPORT PROTEIN SEC7. //1.6e-32:261:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P11075
 F-NT2RM2001659//CARBOXYPEPTIDASE A INHIBITOR. //0.83:30:46//ASCARIS SUUM (PIG ROUNDWORM) (ASCARIS LUMBRICOIDES). //P19399
 F-NT2RM2001664//IKI3 PROTEIN. //1.3e-31:265:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q06706
 F-NT2RM2001668//TONB PROTEIN. //0.32:39:41//XANTHOMONAS CAMPESTRIS (PV. CAMPESTRIS). //034261
 F-NT2RM2001670//ZINC FINGER PROTEIN 174. //3.6e-21:172:39//HOMO SAPIENS (HUMAN). //Q15697
 F-NT2RM2001671//HYPOTHETICAL 118.6 KD PROTEIN C29E 6.03C IN CHROMOSOME I. //1.6e-10:229:24//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09857
 50 F-NT2RM2001675//DIHYDRODIPICOLINATE SYNTHASE (EC

2971

4.2.1.52) (DHDPS). //1.0:184:21//METHANOCOCCUS JANN
ASCHII. //Q57695
F-NT2RM2001681//PROTEIN DISULFIDE ISOMERASE PRECUR
SOR (PDI) (EC 5.3.4.1). //0.0039:199:22//DROSOPHILA
MELANOGASTER (FRUIT FLY). //P54399
F-NT2RM2001688//HYPOTHETICAL 28.1 KD PROTEIN IN SI
PU-PBPC INTERGENIC REGION. //2.6e-21:162:33//BACILL
US SUBTILIS. //P42966
F-NT2RM2001695//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//4.9e-41:60:81//HOMO SAPIENS (HUMAN). //P391 10
94
F-NT2RM2001696//HYPOTHETICAL 24.1 KD PROTEIN IN LE
F4-P33 INTERGENIC REGION. //9.8e-16:126:38//AUTOGRA
PHA CALIFORNICA NUCLEAR POLYHEDROSIS VIRUS (ACMPN
V). //P41479
F-NT2RM2001698//PENAEIDIN-3B PRECURSOR (P3-B). //0.
36:52:34//PENAEUS VANNAMEI (PENOEID SHRIMP) (EUROP
EAN WHITE SHRIMP). //P81059
F-NT2RM2001699//TRANSCRIPTION INITIATION FACTOR TF
IID 30 KD SUBUNIT (TAFII-30) (TAFII30). //0.0012:7 20
9:40//HOMO SAPIENS (HUMAN). //Q12962
F-NT2RM2001700//ACYL-COA DEHYDROGENASE, VERY-LONG-
CHAIN SPECIFIC (EC 1.3.99.-) (VLCAD) (FRAGMENT). //
1.0e-30:140:53//MUS MUSCULUS (MOUSE). //P50544
F-NT2RM2001706//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!//1.5e-33:95:75//HOMO SAPIENS (HUMAN). //P391
95
F-NT2RM2001716//HYPOTHETICAL 118.4 KD PROTEIN IN B
AT2-DAL5 INTERGENIC REGION PRECURSOR. //0.010:116:3
0//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P471 30
79
F-NT2RM2001718//METHYL-ACCEPTING CHEMOTAXIS PROTEI
N TLPB. //0.00029:77:37//BACILLUS SUBTILIS. //P39217
F-NT2RM2001723//POSTERIOR PITUITARY PEPTIDE. //0.9
4:26:53//BOS TAURUS (BOVINE). //P01154
F-NT2RM2001727//E7 PROTEIN. //0.91:46:34//HUMAN PAP
ILLOMAVIRUS TYPE 23. //P50781
F-NT2RM2001730//PROBABLE UBIQUITIN CARBOXYL-TERMIN
AL HYDROLASE K02C4.3 (EC 3.1.2.15) (UBIQUITIN THIO
LESTERASE) (UBIQUITIN-SPECIFIC PROCESSING PROTEAS 40
E) (DEUBIQUITINATING ENZYME). //4.9e-07:139:29//CAE
NORHABDITIS ELEGANS. //Q09931
F-NT2RM2001743//PROENKEPHALIN A PRECURSOR. //0.75:6
5:35//CAVIA PORCELLUS (GUINEA PIG). //P47969
F-NT2RM2001753//HYPOTHETICAL PROTEIN KIAA0210. //1.
5e-14:119:36//HOMO SAPIENS (HUMAN). //Q92609
F-NT2RM2001760//PROTEIN TRANSPORT PROTEIN SEC61 AL
PHA SUBUNIT. //8.3e-58:119:99//CANIS FAMILIARIS (DO
G). //P38377
F-NT2RM2001768//HYPOTHETICAL PROTEIN UL25. //0.45:7 50

2972

7:32//HUMAN CYTOMEGALOVIRUS (STRAIN AD169). //P1676
1
F-NT2RM2001771//ZINC FINGER PROTEIN 135. //4.6e-80:
224:60//HOMO SAPIENS (HUMAN). //P52742
F-NT2RM2001782//MANNOSE-1-PHOSPHATE GUANYLTRANSFER
ASE (EC 2.7.7.13) (ATP-MANNOSE-1-PHOSPHATE GUANYL
YLTRANSFERASE) (NDP-HEXOSE PYROPHOSPHORYLASE). //7.
0e-06:61:45//SACCHAROMYCES CEREVISIAE (BAKER'S YEA
ST). //P41940
F-NT2RM2001784//HYPOTHETICAL PROTEIN UL61. //0.0007
0:145:33//HUMAN CYTOMEGALOVIRUS (STRAIN AD169). //P
16818
F-NT2RM2001785//LINOLEOYL-COA DESATURASE (EC 1.14.
99.25) (DELTA(6)-DESATURASE). //1.5e-08:127:32//SYN
ECHOCYSTIS SP. (STRAIN PCC 6803). //Q08871
F-NT2RM2001797//ZINC FINGER PROTEIN 135. //1.6e-73:
267:49//HOMO SAPIENS (HUMAN). //P52742
F-NT2RM2001800//HYPOTHETICAL HELICASE MG018/MG017/
MG016 HOMOLOG. //3.9e-12:171:33//MYCOPLASMA PNEUMON
IAE. //P75093
F-NT2RM2001803//IKI3 PROTEIN. //1.6e-38:283:36//SAC
CHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q06706
F-NT2RM2001805//COLD SHOCK-LIKE PROTEIN CSPH. //0.5
1:46:32//SALMONELLA TYPHIMURIUM. //Q33793
F-NT2RM2001813//HYPOTHETICAL 40.4 KD TRP-ASP REPEA
TS CONTAINING PROTEIN C14B1.4 IN CHROMOSOME III. //
5.0e-05:82:32//CAENORHABDITIS ELEGANS. //Q17963
F-NT2RM2001823//CHROMODOMAIN-HELICASE-DNA-BINDING
PROTEIN 2 (CHD-2). //3.6e-49:233:45//HOMO SAPIENS
(HUMAN). //Q14647
F-NT2RM2001839//RETICULOCALBIN 1 PRECURSOR. //5.2e-
65:222:56//HOMO SAPIENS (HUMAN). //Q15293
F-NT2RM2001840//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!!//9.6e-33:102:68//HOMO SAPIENS (HUMAN). //P39
194
F-NT2RM2001855//BASP1 PROTEIN. //0.054:120:30//HOMO
SAPIENS (HUMAN). //P80723
F-NT2RM2001867//HYPOTHETICAL 56.6 KD PROTEIN IN UR
E2-SSU72 INTERGENIC REGION. //4.1e-19:88:36//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST). //P53867
F-NT2RM2001879//HYPOTHETICAL 47.3 KD PROTEIN C22C
7.07C IN CHROMOSOME I. //5.9e-15:76:38//SCHIZOSACCH
AROMYCES POMBE (FISSION YEAST). //Q09800
F-NT2RM2001886//HYPOTHETICAL 126.9 KD PROTEIN C22C
7.04 IN CHROMOSOME I. //1.4e-41:249:38//SCHIZOSACCH
AROMYCES POMBE (FISSION YEAST). //Q09798
F-NT2RM2001896//HYPOTHETICAL 83.2 KD PROTEIN IN KA
R4-PBN1 INTERGENIC REGION. //2.1e-59:197:56//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST). //P25582
F-NT2RM2001903//HYPOTHETICAL PROTEIN MJ0263. //0.07

- 0:132:31//METHANOCOCCUS JANNASCHII.//006917
 F-NT2RM2001930//THROMBOSPONDIN 2 PRECURSOR.//7.1e-05:53:47//MUS MUSCULUS(MOUSE).//Q03350
 F-NT2RM2001935//PUTATIVE CUTICLE COLLAGEN F55C10.3.//0.00046:116:35//CAENORHABDITIS ELEGANS.//Q21184
 F-NT2RM2001936//32.3 KD PROTEIN IN CWP1-MBR1 INTERGENIC REGION.//4.5e-27:216:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P28320
 F-NT2RM2001950//HIRUDIN HV1 (BUFRUDIN).//0.59:43:34//HIRUDINARIA MANILLIENSIS (BUFFALO LEECH).//P81492
 F-NT2RM2001982//GUANINE NUCLEOTIDE-BINDING PROTEIN G(1)/G(S)/G(O) GAMMA-8 SUBUNIT (G GAMMA-C).//0.72:35:42//BOS TAURUS (BOVINE).//P50154
 F-NT2RM2001983//PROLINE-RICH PEPTIDE P-B.//0.00035:23:52//HOMO SAPIENS (HUMAN).//P02814
 F-NT2RM2001989//NUCLEOLAR PROTEIN NOP4 (NUCLEOLAR PROTEIN NOP77).//8.6e-24:197:37//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P37838
 F-NT2RM2001997
 F-NT2RM2001998//IMMEDIATE-EARLY PROTEIN IE180.//0.076:92:27//PSEUDORABIES VIRUS (STRAIN INDIANA-FUNK HAUSER / BECKER) (PRV).//P11675
 F-NT2RM2002004//SLF1 PROTEIN.//3.5e-06:235:27//SACCHAROMYCES CEREVISIAE(BAKER'S YEAST).//Q12034
 F-NT2RM2002014//HYPOTHETICAL PROTEIN HI0568.//2.1e-17:235:29//HAEMOPHILUS INFLUENZAE.//P71353
 F-NT2RM2002030//GLUCOSAMINE--FRUCTOSE-6-PHOSPHATE AMINOTRANSFERASE [ISOMERIZING] (EC 2.6.1.16) (HEXOSEPHOSPHATE AMINOTRANSFERASE) (D-FRUCTOSE-6-PHOSPHATE AMIDOTRANSFERASE) (GFAT).//9.5e-105:271:76//MUS MUSCULUS (MOUSE).//P47856
 F-NT2RM2002049//SMALL PROLINE-RICH PROTEIN 2-1.//0.099:41:41//HOMO SAPIENS (HUMAN).//P35326
 F-NT2RM2002055//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS13.//0.012:217:24//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q07878
 F-NT2RM2002088//PUTATIVE HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN X (HNRNP X) (CBP).//1.1e-09:65:53//MUS MUSCULUS (MOUSE).//Q61990
 F-NT2RM2002091//HYPOTHETICAL 57.5 KD PROTEIN IN VM A7-RPS25A INTERGENIC REGION.//0.072:74:37//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53214
 F-NT2RM2002100//ATP-DEPENDENT RNA HELICASE ROK1.//4.5e-50:289:41//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P45818
 F-NT2RM2002109//NT-3 GROWTH FACTOR RECEPTOR PRECURSOR (EC 2.7.1.112) (TRKC TYROSINE KINASE) (GP145-TK) (TRK-C).//1.4e-14:203:32//RATTUS NORVEGICUS (RAT).//Q03351
 F-NT2RM2002128//SKIN SECRETORY PROTEIN XP2 PRECURSOR (APEG PROTEIN).//0.0025:139:31//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P17437
 F-NT2RM2002142//GASTRULATION SPECIFIC PROTEIN G12.//9.2e-20:42:73//BRACHYDANIO RERIO (ZEBRAFISH) (ZEBRA DANIO).//P47805
 F-NT2RM2002145//GLUTENIN, HIGH MOLECULAR WEIGHT SUBUNIT 12 PRECURSOR.//0.0085:200:26//TRITICUM AESTIVUM (WHEAT).//P08488
 F-NT2RM2002178//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//5.8e-05:56:39//BOS TAURUS (BOVINE).//P25508
 F-NT2RM2002580//CCAAT-BINDING TRANSCRIPTION FACTOR SUBUNIT A (CBF-A) (NF-Y PROTEIN CHAIN B) (NF-YB) (CAAT-BOX DNA BINDING PROTEIN SUBUNIT B).//2.9e-14:96:37//PETROMYZON MARINUS (SEA LAMPREY).//P25210
 F-NT2RM4000024//DNA-DIRECTED RNA POLYMERASE III 128 KD POLYPEPTIDE (EC 2.7.7.6) (RNA POLYMERASE III SUBUNIT 2).//8.6e-95:271:67//DROSOPHILA MELANOGASTER (FRUIT FLY).//P25167
 F-NT2RM4000027//INTERFERON-ACTIVATABLE PROTEIN 202 (IFI-202).//0.99:72:31//MUS MUSCULUS (MOUSE).//P15091
 F-NT2RM4000030//LAS1 PROTEIN.//1.4e-14:184:32//SACCHAROMYCES CEREVISIAE(BAKER'S YEAST).//P36146
 F-NT2RM4000046//COLLAGEN ALPHA 1(III) CHAIN (FRAGMENT).//0.99:120:28//RATTUS NORVEGICUS (RAT).//P13941
 F-NT2RM4000061
 F-NT2RM4000085//ATP-DEPENDENT RNA HELICASE A (NUCLEAR DNA HELICASE II) (NDH II) (DEAD BOX PROTEIN 9) (MHEL-5).//8.5e-40:263:38//MUS MUSCULUS (MOUSE).//070133
 F-NT2RM4000086//HYPOTHETICAL PROTEIN HI1497.//1.0:27:37//HAEMOPHILUS INFLUENZAE.//P44221
 F-NT2RM4000104//ZINC FINGER PROTEIN 134.//1.0e-26:64:56//HOMO SAPIENS (HUMAN).//P52741
 F-NT2RM4000139//PREPROTEIN TRANSLOCASE SECE SUBUNIT.//0.99:38:42//THERMOTOGA MARITIMA.//P35874
 F-NT2RM4000155//THREONYL-TRNA SYNTHETASE, CYTOPLASMIC (EC 6.1.1.3) (THREONINE--TRNA LIGASE) (THRRS).//6.3e-34:181:40//HOMO SAPIENS (HUMAN).//P26639
 F-NT2RM4000156//EXTENSIN PRECURSOR (CELL WALL HYDROXYPROLINE-RICH GLYCOPROTEIN).//4.6e-12:142:33//NICOTIANA TABACUM (COMMON TOBACCO).//P13983
 F-NT2RM4000167//KINESIN-LIKE PROTEIN KIF4.//3.4e-123:269:91//MUS MUSCULUS (MOUSE).//P33174
 F-NT2RM4000169//M PROTEIN, SEROTYPE 2.2 PRECURSOR.//9.7e-10:229:26//STREPTOCOCCUS PYOGENES.//P50469

2975

F-NT2RM4000191//P68-LIKE PROTEIN.//2.1e-11:104:40/
/SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P24783
F-NT2RM4000197//CUTICLE PROTEIN CP463 (CPCP463).//
0.84:29:37//CANCER PAGURUS (ROCK CRAB).//P81587
F-NT2RM4000199//SALIVARY PROLINE-RICH PROTEIN PO
(ALLELE K) [CONTAINS: PEPTIDE P-D] (FRAGMENT).//1.
8e-06:187:34//HOMO SAPIENS (HUMAN).//P10162
F-NT2RM4000200//HYPOTHETICAL 9.4 KD PROTEIN IN FLA
L 3' REGION (ORF3).//0.52:42:40//BACILLUS LICHENIFO
RMIS.//P22754
F-NT2RM4000202//COLLAGEN ALPHA 1(VI11) CHAIN PRECU
RSOR (ENDOTHELIAL COLLAGEN).//0.00044:168:32//ORYC
TOLAGUS CUNICULUS (RABBIT).//P14282
F-NT2RM4000210//EXTENSIN PRECURSOR.//0.27:129:27//
DAUCUS CAROTA (CARROT).//P06599
F-NT2RM4000215//MAK16 PROTEIN.//2.0e-65:234:52//SA
CCHAROMYCES CEREVISIAE(BAKER'S YEAST).//P10962
F-NT2RM4000229//GAR2 PROTEIN.//0.13:217:26//SCHIZO
SACCHAROMYCES POMBE (FISSION YEAST).//P41891
F-NT2RM4000233//TRANSMEMBRANE PROTEIN SEX PRECURSO
R.//0.047:108:30//HOMOSAPIENS (HUMAN).//P51805
F-NT2RM4000244//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//0.67:59:27//BALAENOPTERA PHYSALUS (FINB
ACK WHALE) (COMMON RORQUAL).//P24947
F-NT2RM4000251//PROLINE-RICH PROTEIN MP-3 (FRAGMEN
T).//0.0059:108:35//MUS MUSCULUS (MOUSE).//P05143
F-NT2RM4000265//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//8.1e-38:70:70//HOMO SAPIENS (HUMAN).//P3918
8
F-NT2RM4000290//TRANSDUCIN-LIKE ENHANCER PROTEIN 3
(ESG3).//1.6e-115:209:94//HOMO SAPIENS (HUMAN).//
Q04726
F-NT2RM4000324//PRESPORE PROTEIN DP87 PRECURSOR.//
0.14:136:30//DICTYOSTELIUM DISCOIDEUM (SLIME MOL
D).//Q04503
F-NT2RM4000327//HYPOTHETICAL 8.9 KD PROTEIN IN IE0
-IE1 INTERGENIC REGION.//0.91:73:28//AUTOGRAPHICA CA
LIFORNICA NUCLEAR POLYHEDROSIS VIRUS (ACMPV).//P4
1703
F-NT2RM4000344//YME1 PROTEIN HOMOLOG (EC 3.4.24.
-).//9.4e-78:241:55//CAENORHABDITIS ELEGANS.//P548
13
F-NT2RM4000349//CYSTEINE STRING PROTEIN (CCCS1).//
0.055:22:59//TORPEDO CALIFORNICA (PACIFIC ELECTRIC
RAY).//P56101
F-NT2RM4000354//LETHAL(2)DENTICLELESS PROTEIN (DTL
83 PROTEIN).//4.6e-26:208:35//DROSOPHILA MELANOGAS
TER (FRUIT FLY).//Q24371
F-NT2RM4000356//COAT PROTEIN.//0.11:105:36//SATELL
ITE TOBACCO MOSAIC VIRUS (STMV).//P17574

2976

F-NT2RM4000366//IMMEDIATE-EARLY PROTEIN.//1.2e-05:
215:24//HERPESVIRUS SAIMIRI (STRAIN 11).//Q01042
F-NT2RM4000368//HYPOTHETICAL 7.3 KD PROTEIN IN RPB
A-GP46 INTERGENIC REGION.//0.54:46:36//BACTERIOPHA
GE RB69.//064300
F-NT2RM4000386//RHSC PROTEIN PRECURSOR.//0.0096:16
2:29//ESCHERICHIA COLI.//P16918
F-NT2RM4000395//HYPOTHETICAL 52.9 KD PROTEIN IN SA
P155-YMR31 INTERGENIC REGION.//4.5e-66:256:53//SACC
HAROMYCES CEREVISIAE (BAKER'S YEAST).//P43616
F-NT2RM4000414//HYPOTHETICAL 6.0 KD PROTEIN IN THI
12 5' REGION.//0.13:33:48//SACCHAROMYCES CEREVISIAE
(BAKER'S YEAST).//P53820
F-NT2RM4000421//MRNA TRANSPORT REGULATOR MTR10.//
5.0e-13:171:29//SACCHAROMYCES CEREVISIAE (BAKER'S
YEAST).//Q99189
F-NT2RM4000425//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!//2.1e-25:46:80//HOMO SAPIENS (HUMAN).//P391
93
F-NT2RM4000433//CUTICLE COLLAGEN 3A3.//2.5e-06:77:
38//HAEMONCHUS CONTORTUS.//P16253
F-NT2RM4000457//HYPOTHETICAL 111.9 KD PROTEIN C22H
10.03C IN CHROMOSOME I.//4.3e-09:215:22//SCHIZOSAC
CHAROMYCES POMBE (FISSION YEAST).//Q10297
F-NT2RM4000471//TRNA SPLICING PROTEIN SPL1.//6.7e-
73:163:65//CANDIDA ALBICANS (YEAST).//P87185
F-NT2RM4000486//COLLAGEN ALPHA 2(VI) CHAIN PRECURS
OR.//0.0012:121:34//GALLUS GALLUS (CHICKEN).//P159
88
F-NT2RM4000496//DNA-DIRECTED RNA POLYMERASE II LAR
GEST SUBUNIT (EC 2.7.7.6) (RNA POLYMERASE II SUBUN
IT 1).//5.9e-09:175:35//SCHIZOSACCHAROMYCESPOMBE
(FISSION YEAST).//P36594
F-NT2RM4000511//SALIVARY GLUE PROTEIN SGS-3 PRECUR
SOR.//0.020:122:31//DROSOPHILA SIMULANS (FRUIT FL
Y).//P13729
F-NT2RM4000514//ATP SYNTHASE A CHAIN (EC 3.6.1.34)
(PROTEIN 6).//0.46:68:32//ARTEMIA SANFRANCISCANA
(BRINE SHRIMP) (ARTEMIA FRANCISCANA).//Q37708
F-NT2RM4000515//GAR2 PROTEIN.//3.2e-05:198:27//SCH
IZOSACCHAROMYCES POMBE(FISSION YEAST).//P41891
F-NT2RM4000520//HYPOTHETICAL 7.5 KD PROTEIN (ORF 6
3).//0.011:55:38//SPINACIA OLERACEA (SPINACH).//PO
8974
F-NT2RM4000531//ZINC FINGER PROTEIN 169 (FRAGMEN
T).//3.6e-44:244:42//HOMO SAPIENS (HUMAN).//Q14929
F-NT2RM4000532//PUTATIVE MEMBRANE PROTEIN 53.//1.
0:47:34//HERPESVIRUS SAIMIRI (STRAIN 11).//Q01049
F-NT2RM4000534//HYPOTHETICAL 5.9 KD PROTEIN IN WRB
A-PUTA INTERGENIC REGION.//0.75:26:46//ESCHERICHIA

- COLI.//P56614
 F-NT2RM4000585//CAG POLYPROTEIN [CONTAINS: CORE PROTEIN P16; CORE PROTEIN P26].//0.019:86:34//HUMAN IMMUNODEFICIENCY VIRUS TYPE 2 (ISOLATE SBLISY) (HIV-2).//P12450
 F-NT2RM4000590//RING CANAL PROTEIN (KELCH PROTEIN).//5.0e-23:224:29//DROSOPHILA MELANOCASTER (FRUIT FLY).//Q04652
 F-NT2RM4000595//HYPOTHETICAL 54.9 KD PROTEIN C02F 5.7 IN CHROMOSOME III.//3.8e-62:226:50//CAENORHABDITIS ELEGANS.//P34284
 F-NT2RM4000603//SRC SUBSTRATE CORTACTIN (AMPLAXIN) (EMS1 ONCOGENE).//0.077:132:22//HOMO SAPIENS (HUMAN).//Q14247
 F-NT2RM4000611//HYPOTHETICAL 40.4 KD TRP-ASP REPEATS CONTAINING PROTEIN C14B1.4 IN CHROMOSOME III.//1.9e-06:82:32//CAENORHABDITIS ELEGANS.//Q17963
 F-NT2RM4000616//ACETYL-COENZYME A SYNTHETASE (EC 6.2.1.1) (ACETATE--COALIGASE) (ACYL- ACTIVATING ENZYME).//5.3e-79:213:62//ESCHERICHIA COLI.//P27550
 F-NT2RM4000674//HYPOTHETICAL SYMPORTER SLL1374.//1.3e-11:147:32//SYNECHOCYSTIS SP. (STRAIN PCC 6803).//P74168
 F-NT2RM4000689
 F-NT2RM4000698//CHORION CLASS HIGH-CYSTEINE HCA PROTEIN 12 PRECURSOR (HC-A.12).//0.26:45:33//BOMBYX MORI (SILK MOTH).//P05687
 F-NT2RM4000700//THIOPHENE AND FURAN OXIDATION PROTEIN THDF.//0.95:165:25//BORRELIA BURGDORFERI (LYME DISEASE SPIROCHETE).//P53364
 F-NT2RM4000712//PROBABLE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE R10E11.3(EC 3.1.2.15) (UBIQUITIN THIOLESTERASE) (UBIQUITIN-SPECIFIC PROCESSING PROTEASE) (DEUBIQUITINATING ENZYME).//2.2e-82:152:63//CAENORHABDITIS ELEGANS.//P34547
 F-NT2RM4000717//SALIVARY GLUE PROTEIN SCS-3 PRECURSOR.//0.80:54:40//DROSOPHILA SIMULANS (FRUIT FLY).//P13729
 F-NT2RM4000733//OCTAPEPTIDE-REPEAT PROTEIN T2.//1.5e-08:139:28//MUS MUSCULUS (MOUSE).//Q06666
 F-NT2RM4000734//GASTRULA ZINC FINGER PROTEIN XLCCF 26.1 (FRAGMENT).//7.2e-20:205:28//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P18715
 F-NT2RM4000741//SPERM PROTAMINE P1.//0.89:52:38//ISOODON MACRORUS (SHORT-NOSED BANDICOOT).//P42136
 F-NT2RM4000751//ZINC FINGER PROTEIN 26 (ZFP-26) (MKR3 PROTEIN) (FRAGMENT).//5.2e-77:246:52//MUS MUSCULUS (MOUSE).//P10076
 F-NT2RM4000764//KERATIN, GLYCINE/TYROSINE-RICH OF HAIR.//0.062:33:42//OVIS ARIES (SHEEP).//Q02958
 F-NT2RM4000778
 F-NT2RM4000779//SULFATED SURFACE GLYCOPROTEIN 185 (SSG 185).//0.014:53:45//VOLVOX CARTERI.//P21997
 F-NT2RM4000787//BONE MORPHOGENETIC PROTEIN 1 PRECURSOR (EC 3.4.24.-) (BMP-1).//0.00011:73:39//MUS MUSCULUS (MOUSE).//P98063
 F-NT2RM4000790//SPORE COAT PROTEIN SP96.//0.00083:157:29//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P14328
 F-NT2RM4000795//CHOLINESTERASE PRECURSOR (EC 3.1.1.8) (ACYLCHOLINE ACYLHYDROLASE) (CHOLINE ESTERASE II) (BUTYRYLCHOLINE ESTERASE) (PSEUDOCOLINESTERASE).//7.4e-41:271:36//HOMO SAPIENS (HUMAN).//P06276
 F-NT2RM4000796//5-METHYLCYTOSINE-SPECIFIC RESTRICTION ENZYME B (EC 3.1.21.-).//0.28:82:30//ESCHERICHIA COLI.//P15005
 F-NT2RM4000798//PROTEIN TRANSPORT PROTEIN SEC7.//4.7e-38:165:48//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P11075
 F-NT2RM4000813//METALLOTHIONEIN-IB.//0.0025:25:44//OVIS ARIES (SHEEP).//P09577
 F-NT2RM4000820
 F-NT2RM4000833//HYPOTHETICAL PROTEIN MJ1136.//6.5e-42:206:41//METHANOCOCCUS JANNASCHII.//Q58536
 F-NT2RM4000848//BRAIN-SPECIFIC HOMEBOX/POU DOMAIN PROTEIN 3A (BRN-3A) (BRN-3.0).//0.00060:159:33//MUS MUSCULUS (MOUSE).//P17208
 F-NT2RM4000852//SMALL PROLINE-RICH PROTEIN 2B (SPR-2B).//0.0076:13:69//HOMO SAPIENS (HUMAN).//P35325
 F-NT2RM4000855//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//0.0060:68:44//HOMO SAPIENS (HUMAN).//P39194
 F-NT2RM4000887//RTS1 PROTEIN (SCS1 PROTEIN).//0.23:153:24//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38903
 F-NT2RM4000895//HYPOTHETICAL 53.5 KD PROTEIN IN PH02-POL3 INTERGENIC REGION.//3.3e-09:80:46//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P43123
 F-NT2RM4000950//HYPOTHETICAL PROTEIN MJ0572.//0.090:68:29//METHANOCOCCUS JANNASCHII.//Q57992
 F-NT2RM4000971//KINESIN LIGHT CHAIN (KLC).//0.79:201:24//LOLIGO PEALEII (LONGFIN SQUID).//P46825
 F-NT2RM4000979//MYOSIN REGULATORY LIGHT CHAIN 2, NONSARCOMERIC (MYOSIN RLC).//1.2e-07:25:96//HOMO SAPIENS (HUMAN).//P19105
 F-NT2RM4000996//ZINC FINGER PROTEIN 37 (ZFP-37) (MALE GERM CELL SPECIFIC ZINC FINGER PROTEIN).//1.4e-56:253:46//MUS MUSCULUS (MOUSE).//P17141
 F-NT2RM4001002

F-NT2RM4001016//GAG POLYPROTEIN [CONTAINS: CORE PROTEIN P15; INNER COATPROTEIN P12; CORE SHELL PROTEIN P30].//0.25:101:31//FBR MURINE OSTEOSARCOMA VIRUS.//P29175

F-NT2RM4001032//CUTICLE COLLAGEN 2.//2.6e-07:130:39//CAENORHABDITIS ELEGANS.//P17656

F-NT2RM4001047//MO25 PROTEIN.//5.6e-107:252:80//MUS MUSCULUS (MOUSE).//Q06138

F-NT2RM4001054//PROTEIN TRANSPORT PROTEIN SEC61 ALPHA SUBUNIT.//9.0e-109:209:94//CANIS FAMILIARIS (DOG).//P38377

F-NT2RM4001084//HYPOTHETICAL TRANSCRIPTIONAL REGULATOR IN UXUR-IADA INTERGENIC REGION.//0.57:95:30//ESCHERICHIA COLI.//P39376

F-NT2RM4001092//HYPOTHETICAL 127.4 KD PROTEIN F07F6.4 IN CHROMOSOME III.//2.5e-47:231:47//CAENORHABDITIS ELEGANS.//Q09531

F-NT2RM4001116//HYPOTHETICAL 216.3 KD PROTEIN R06F6.8 IN CHROMOSOME II.//1.3e-08:243:23//CAENORHABDITIS ELEGANS.//Q09417

F-NT2RM4001140//HOMEODOMAIN PROTEIN MSH-D.//7.1e-13:103:38//BRACHYDANI0 RERIO (ZEBRAFISH) (ZEBRA DANI0).//Q01704

F-NT2RM4001151//SYNAPSINS IA AND IB (BRAIN PROTEIN 4.1).//0.26:96:34//HOMO SAPIENS (HUMAN).//P17600

F-NT2RM4001155//ADRENAL MEDULLA 50 KD PROTEIN.//3.6e-103:201:91//BOS TAURUS (BOVINE).//Q27969

F-NT2RM4001160//GLUTATHIONE S-TRANSFERASE (EC 2.5.1.18) (CLASS-PHI) (FRAGMENTS).//1.0:33:36//BRASSICA OLERACEA (CAULIFLOWER).//P48438

F-NT2RM4001187//PREPROTEIN TRANSLOCASE SECA SUBUNIT.//0.44:158:27//MYCOPLASMA GENITALIUM.//P47318

F-NT2RM4001191//LONG NEUROTOXIN 2 (TOXIN C).//0.99:44:43//ASTROTIA STOKESI (STOKES' S SEA SNAKE) (DI STEIRA STOKESI).//P01381

F-NT2RM4001200//ZINC FINGER PROTEIN 135.//2.2e-82:245:59//HOMO SAPIENS (HUMAN).//P52742

F-NT2RM4001203//HYPOTHETICAL 57.5 KD PROTEIN IN VM A7-RPS25A INTERGENIC REGION.//0.028:94:40//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53214

F-NT2RM4001204//SPLICIOSOME ASSOCIATED PROTEIN 62 (SAP 62) (SF3A66).//0.0096:182:34//HOMO SAPIENS (HUMAN).//Q15428

F-NT2RM4001217//RING CANAL PROTEIN (KELCH PROTEIN).//2.1e-21:221:29//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q04652

F-NT2RM4001256//CBP3 PROTEIN PRECURSOR.//0.30:55:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P21560

F-NT2RM4001258//HYPOTHETICAL 35.5 KD PROTEIN IN TR 50

ANSPOSON TN4556.//0.00031:132:39//STREPTOMYCES FRATIAE.//P20186

F-NT2RM4001309//SALIVARY PROLINE-RICH PROTEIN PRECURSOR (CLONE CP7) [CONTAINS: BASIC PEPTIDE P-F] (FRAGMENT).//0.048:132:28//HOMO SAPIENS (HUMAN).//P02812

F-NT2RM4001313//PHOSPHATIDYLINOSITOL 3-KINASE VPS34-LIKE (EC 2.7.1.137) (PI3-KINASE) (PTDINS-3-KINASE) (PI3K).//2.6e-37:124:65//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P54676

F-NT2RM4001316//ACYL-COA DEHYDROGENASE, MEDIUM-CHAIN SPECIFIC PRECURSOR (EC 1.3.99.3) (MCAD).//1.7e-10:185:30//RATTUS NORVEGICUS (RAT).//P08503

F-NT2RM4001320//PUTATIVE RHO/RAC GUANINE NUCLEOTIDE EXCHANGE FACTOR (RHO/RAC GEF) (FACIOGENITAL DYSPLASIA PROTEIN HOMOLOG).//1.5e-08:197:26//MUS MUSCULUS (MOUSE).//P52734

F-NT2RM4001340//UTR4 PROTEIN (UNKNOWN TRANSCRIPT 4 PROTEIN).//7.7e-14:82:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P32626

F-NT2RM4001344//HYPOTHETICAL GTP-BINDING PROTEIN IN POP2-HOL1 INTERGENIC REGION.//3.3e-16:128:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53742

F-NT2RM4001347//HYPOTHETICAL 76.9 KD PROTEIN IN RP M2-TUB1 INTERGENIC REGION.//0.067:111:33//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q04511

F-NT2RM4001371

F-NT2RM4001382//HISTIDINE-RICH GLYCOPROTEIN PRECURSOR.//1.0e-08:82:39//PLASMODIUM LOPHURAE.//P04929

30 F-NT2RM4001384

F-NT2RM4001410//A-AGGLUTININ ATTACHMENT SUBUNIT PRECURSOR.//2.1e-08:185:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P32323

F-NT2RM4001411//EARLY NODULIN 20 PRECURSOR (N-20).//5.3e-05:105:38//MEDICAGO TRUNCATULA (BARREL MEDIC).//P93329

F-NT2RM4001412//GTPASE-ACTIVATING PROTEIN (GAP) (RAS P21 PROTEIN ACTIVATOR) (P120GAP) (RASGAP).//6.2e-17:109:41//RATTUS NORVEGICUS (RAT).//P50904

40 F-NT2RM4001414//ZINC FINGER PROTEIN 177.//8.3e-06:54:50//HOMO SAPIENS (HUMAN).//Q13360

F-NT2RM4001437//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!//2.1e-24:87:65//HOMO SAPIENS (HUMAN).//P39192

F-NT2RM4001444//PROBABLE ISOLEUCYL-TRNA SYNTHETASE (EC 6.1.1.5) (ISOLEUCINE--TRNA LIGASE) (ILERS) (FRAGMENT).//2.6e-45:197:47//CIONA INTESTINALIS.//Q94425

F-NT2RM4001454//HYPOTHETICAL PROTEIN KIAA0041 (FRAGMENT).//0.0060:95:29//HOMO SAPIENS (HUMAN).//Q150

57

F-NT2RM4001455//PROBABLE E5B PROTEIN.//0.41:44:36/
/HUMAN PAPILOMAVIRUS TYPE 6B.//P06461

F-NT2RM4001483//ZINC FINGER PROTEIN 136.//1.7e-28:
85:64//HOMO SAPIENS (HUMAN).//P52737

F-NT2RM4001489//PTB-ASSOCIATED SPLICING FACTOR (PS
F).//0.086:111:34//HOMO SAPIENS (HUMAN).//P23246

F-NT2RM4001519//ACID UREASE ALPHA SUBUNIT (EC 3.5.
1.5) (UREA AMIDOHYDROLASE).//0.82:51:47//LACTOBACI
LLUS FERMENTUM.//P26929

F-NT2RM4001522//TROPOMYOSIN.//0.030:117:23//SCHIZO
SACCHAROMYCES POMBE (FISSION YEAST).//Q02088

F-NT2RM4001557

F-NT2RM4001565//HYPOTHETICAL 44.3 KD PROTEIN C1F7.
07C IN CHROMOSOME I.//0.99:42:40//SCHIZOSACCHAROMY
CES POMBE (FISSION YEAST).//Q09919

F-NT2RM4001566//GLUCOAMYLASE S1/S2 PRECURSOR (EC
3.2.1.3) (GLUCAN 1,4-ALPHA- GLUCOSIDASE) (1,4-ALPH
A-D-GLUCAN GLUCOHYDROLASE).//0.054:190:23//SACCHAR
OMYCES CEREVISIAE (BAKER'S YEAST).//P08640

F-NT2RM4001569//GUANINE NUCLEOTIDE-BINDING PROTEIN
BETA SUBUNIT-LIKE PROTEIN (ACTIVATED PROTEIN KINA
SE C RECEPTOR HOMOLOG).//0.72:64:31//TRYPANOSOMA B
RUCI BRUCEI.//Q94775

F-NT2RM4001582

F-NT2RM4001592//DNA REPAIR PROTEIN RAD9.//0.00037:
198:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//
P14737

F-NT2RM4001594//IMMEDIATE-EARLY PROTEIN IE180.//1.
9e-05:147:34//PSEUDORABIES VIRUS (STRAIN KAPLAN)
(PRV).//P33479

F-NT2RM4001597//THIOL: DISULFIDE INTERCHANGE PROTEI
N TLPA (CYTOCHROME C BIOGENESIS PROTEIN TLPA).//5.
7e-06:122:29//BRADYRHIZOBIUM JAPONICUM.//P43221

F-NT2RM4001605//NUCLEAR PORE COMPLEX PROTEIN NUP15
5 (NUCLEOPORIN NUP155) (155 KD NUCLEOPORIN) (P140).
//1.7e-128:249:96//RATTUS NORVEGICUS (RAT).//P3719
9

F-NT2RM4001611//SIS2 PROTEIN (HALOTOLERANCE PROTEI
N HAL3).//1.5e-35:128:47//SACCHAROMYCES CEREVISIAE
(BAKER'S YEAST).//P36024

F-NT2RM4001629//MAGUK P55 SUBFAMILY MEMBER 3 (MPP3
PROTEIN) (DISCS, LARGE HOMOLOG 3).//5.8e-42:254:3
7//HOMO SAPIENS (HUMAN).//Q13368

F-NT2RM4001650//HOMEODOMAIN PROTEIN HOX-A4 (CHOX-1.
4).//0.62:19:57//GALLUS GALLUS (CHICKEN).//P17277

F-NT2RM4001662//PROTEIN KINASE C, ALPHA TYPE (EC
2.7.1.-) (PKC-ALPHA).//0.29:90:32//HOMO SAPIENS (H
UMAN).//P17252

F-NT2RM4001666//HYPOTHETICAL 48.6 KD PROTEIN IN AL 50

PA-GABP INTERGENIC REGION.//1.1e-31:137:44//ESCHER
ICHIA COLI.//P37339

F-NT2RM4001682//PROBABLE 60S RIBOSOMAL PROTEIN L2
2.//0.98:55:29//CAENORHABDITIS ELEGANS.//P52819

F-NT2RM4001710//HYPOTHETICAL PROTEIN KIAA0039 (FRA
GMENT).//0.56:113:28//HOMO SAPIENS (HUMAN).//Q1505
4

F-NT2RM4001714//SEPTIN 2 HOMOLOG (FRAGMENT).//1.4e
-108:255:77//HOMO SAPIENS (HUMAN).//Q14141

10 F-NT2RM4001715//HYPOTHETICAL PROTEIN C19G10.16 IN
CHROMOSOME I (FRAGMENT).//2.1e-36:148:38//SCHIZOSA
CCHAROMYCES POMBE (FISSION YEAST).//Q10342

F-NT2RM4001731//HYPOTHETICAL 54.9 KD PROTEIN C02F
5.7 IN CHROMOSOME III.//1.1e-05:90:33//CAENORHABDI
TIS ELEGANS.//P34284

F-NT2RM4001741//TALIN.//1.1e-106:208:99//MUS MUSCU
LUS (MOUSE).//P26039

F-NT2RM4001746//EBNA-1 NUCLEAR PROTEIN.//1.6e-09:1
55:38//EPSTEIN-BARR VIRUS (STRAIN B95-8) (HUMAN HE
RPESVIRUS 4).//P03211

F-NT2RM4001754//COLLAGEN ALPHA 5(IV) CHAIN PRECURS
OR.//0.93:158:33//HOMOSAPIENS (HUMAN).//P29400

F-NT2RM4001758//PUTATIVE SERINE/THREONINE-PROTEIN
KINASE P78 (EC 2.7.1.-).//5.1e-113:277:79//HOMO SA
PIENS (HUMAN).//P27448

F-NT2RM4001776//MYOSIN I ALPHA (MMI-ALPHA).//2.2e-
73:262:54//MUS MUSCULUS (MOUSE).//P46735

F-NT2RM4001783//ZINC FINGER PROTEIN HRX (ALL-1) (F
RAGMENT).//5.3e-26:169:39//MUS MUSCULUS (MOUSE).//
P55200

F-NT2RM4001810//MALE SPECIFIC SPERM PROTEIN MST84D
B.//2.3e-05:68:42//DROSOPHILA MELANOGASTER (FRUIT
FLY).//Q01643

F-NT2RM4001813//RHODOCETIN ALPHA SUBUNIT.//2.3e-0
5:115:34//ACKISTRODON RHODOSTOMA (MALAYAN PIT VIPE
R) (CALLOSELASMA RHODOSTOMA).//P81397

F-NT2RM4001819//CELL SURFACE GLYCOPROTEIN EMR1 PRE
CURSOR (EMR1 HORMONE RECEPTOR) (CELL SURFACE GLYCO
PROTEIN F4/80).//1.7e-06:159:25//MUS MUSCULUS (MOU
SE).//Q61549

F-NT2RM4001823//ZINC FINGER PROTEIN ZIC1 (ZINC FIN
GER PROTEIN OF THE CEREBELLUM 1).//2.6e-18:114:40/
/MUS MUSCULUS (MOUSE).//P46684

F-NT2RM4001828//ZINC FINGER PROTEIN 84 (ZINC FINGE
R PROTEIN HPF2).//4.0e-81:253:59//HOMO SAPIENS (HU
MAN).//P51523

F-NT2RM4001836//SPIDROIN 2 (DRAGLINE SILK FIBROIN
2) (FRAGMENT).//0.21:176:30//NEPHILA CLAVIPES (ORB
SPIDER).//P46804

F-NT2RM4001841//PROLINE-RICH PEPTIDE P-B.//0.046:2

2983

2984

- 7:40//HOMO SAPIENS (HUMAN).//P02814
 F-NT2RM4001842//HYPOTHETICAL 7.0 KD PROTEIN B03B8.
 1 IN CHROMOSOME III.//0.98:35:42//CAENORHABDITIS E
 LEGANS.//Q11104
 F-NT2RM4001856//HYPOTHETICAL 75.2 KD PROTEIN IN AC
 S1-GCV3 INTERGENIC REGION.//2.3e-37:242:37//SACCHA
 ROMYCES CEREVISIAE (BAKER'S YEAST).//P39722
 F-NT2RM4001858//T-BOX PROTEIN VECT (T-BOX PROTEIN
 BRAT) (T-BOX PROTEIN ANTIPODEAN).//1.8e-23:78:64//
 XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P87377
 F-NT2RM4001865//NEURONAL CALCIUM SENSOR 2 (NCS-2).
 //0.012:83:28//CAENORHABDITIS ELEGANS.//P36609
 F-NT2RM4001876//HYPOTHETICAL 118.4 KD PROTEIN IN B
 AT2-DAL5 INTERGENIC REGION PRECURSOR.//3.8e-10:24
 2:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P
 47179
 F-NT2RM4001880//EC PROTEIN HOMOLOG.//0.22:59:32//A
 RABIDOPSIS THALIANA (MOUSE-EAR CRESS).//P93746
 F-NT2RM4001905//60S RIBOSOMAL PROTEIN L40 (CEP52).
 //0.57:20:60//HOMO SAPIENS (HUMAN), RATTUS NORVEGI
 CUS (RAT), AND GALLUS GALLUS (CHICKEN).//P14793
 F-NT2RM4001922
 F-NT2RM4001930//PUTATIVE GLUCOSYLTRANSFERASE C08B1
 1.8 (EC 2.4.1.-).//5.5e-45:167:53//CAENORHABDITIS
 ELEGANS.//Q09226
 F-NT2RM4001938//RTOA PROTEIN (RATIO-A).//0.0036:12
 0:32//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P546
 81
 F-NT2RM4001940//IROQUOIS-CLASS HOMEODOMAIN PROTEIN
 IRX-1 (FRAGMENT).//0.32:31:48//HOMO SAPIENS (HUMA
 N).//P78415
 F-NT2RM4001953//!!!! ALU SUBFAMILY SC WARNING ENTR
 Y !!!!!//2.2e-43:56:85//HOMO SAPIENS (HUMAN).//P391
 92
 F-NT2RM4001965//IG ALPHA-1 CHAIN C REGION.//0.56:7
 3:34//GORILLA GORILLAGORILLA (LOWLAND GORILLA).//P
 20758
 F-NT2RM4001969//SALIVARY PROLINE-RICH PROTEIN PREC
 URSOR (CLONES CP3, CP4AND CP5) [CONTAINS: BASIC PE
 PTIDE IB-6; PEPTIDE P-H].//0.0016:140:27//HOMO SAP
 IENS (HUMAN).//P04280
 F-NT2RM4001979//ZINC FINGER PROTEIN 84 (ZINC FINGE
 R PROTEIN HPF2).//3.9e-21:103:51//HOMO SAPIENS (HU
 MAN).//P51523
 F-NT2RM4001984//HYPOTHETICAL PROTEIN LAMBDA-SP5.//
 0.0034:50:40//MUS MUSCULUS (MOUSE).//P15974
 F-NT2RM4001987//IRREGULAR CHIASM C-ROUGHEST PROTEI
 N PRECURSOR (IRREC PROTEIN).//6.9e-17:115:31//DROS
 OPHILA MELANOGASTER (FRUIT FLY).//Q08180
 F-NT2RM4002013//HYPOTHETICAL 54.5 KD TRP-ASP REPEA
 50
 TS CONTAINING PROTEINZC302.2 IN CHROMOSOME V.//0.0
 062:117:28//CAENORHABDITIS ELEGANS.//Q23256
 F-NT2RM4002018//SPORE COAT PROTEIN SP96.//4.3e-06:
 203:28//DICTYOSTELIUMDISCOIDEUM (SLIME MOLD).//P14
 328
 F-NT2RM4002034//RHO-GAP HEMATOPOIETIC PROTEIN C1
 (P115) (KIAA0131).//0.78:132:25//HOMO SAPIENS (HUM
 AN).//P98171
 F-NT2RM4002044//VITELLOGENIN I PRECURSOR (MINOR VI
 TELLOGENIN) [CONTAINS: LIPOVITELLIN I (LVI); PHOSVI
 TIN (PV); LIPOVITELLIN II (LVI1); YGP42].//0.062:2
 01:24//GALLUS GALLUS (CHICKEN).//P87498
 F-NT2RM4002054//DUPLICATE PROCYCLIN.//0.0079:44:52
 //TRYPANOSOMA BRUCEI BRUCEI.//P14044
 F-NT2RM4002055//PUTATIVE Z PROTEIN.//0.82:39:30//O
 VIS ARIES (SHEEP).//P08105
 F-NT2RM4002062//ASPARTYL-TRNA SYNTHETASE (EC 6.1.
 1.12) (ASPARTATE--TRNALIGASE) (ASPRS).//7.0e-37:8
 0:52//THERMUS AQUATICUS (SUBSP. THERMOPHILUS).//P3
 6419
 F-NT2RM4002063//SARCOSINE OXIDASE (EC 1.5.3.1).//
 2.2e-25:216:31//BACILLUS SP. (STRAIN NS-129).//P23
 342
 F-NT2RM4002066//HYPOTHETICAL PROTEIN KIAA0192 (FRA
 GMENT).//1.1e-94:260:71//HOMO SAPIENS (HUMAN).//Q9
 3074
 F-NT2RM4002067//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!//1.5e-15:51:70//HOMO SAPIENS (HUMAN).//P3918
 8
 F-NT2RM4002073//ELASTIN PRECURSOR (TROPOELASTIN)./
 /4.9e-05:88:36//HOMO SAPIENS (HUMAN).//P15502
 F-NT2RM4002075//RING CANAL PROTEIN (KELCH PROTEI
 N).//7.2e-43:220:41//DROSOPHILA MELANOGASTER (FRUI
 T FLY).//Q04652
 F-NT2RM4002093//POLYPYRIMIDINE TRACT-BINDING PROTE
 IN (PTB) (HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN
 I) (HNRNP I) (57 KD RNA-BINDING PROTEIN PPTB-1).//
 1.8e-93:255:72//HOMO SAPIENS (HUMAN).//P26599
 F-NT2RM4002109//KINESIN-LIKE PROTEIN KIF4.//3.7e-1
 01:260:78//MUS MUSCULUS (MOUSE).//P33174
 F-NT2RM4002128//HYPOTHETICAL PROTEIN IN CYCB 3' REG
 ION PRECURSOR (ORF2) (FRAGMENT).//0.91:49:32//PARA
 COCCUS DENITRIFICANS.//P29969
 F-NT2RM4002140//GROUCHO PROTEIN (ENHANCER OF SPLIT
 M9/10).//0.36:104:22//DROSOPHILA MELANOGASTER (FR
 UIT FLY).//P16371
 F-NT2RM4002145//SLIT PROTEIN PRECURSOR.//8.6e-13:1
 27:33//DROSOPHILA MELANOGASTER (FRUIT FLY).//P2401
 4
 F-NT2RM4002146//MAGO NASHI PROTEIN.//7.9e-69:143:9

2985

1//DROSOPHILA MELANOGASTER (FRUIT FLY).//P49028
 F-NT2RM4002161//DUAL SPECIFICITY PROTEIN PHOSPHATASE (EC 3.1.3.48) (EC 3.1.3.16).//0.0062:99:26//CHLAMYDOMONAS EUCAMETOS.//Q39491
 F-NT2RM4002174//MRP PROTEIN.//4.5e-50:183:55//ESCHERICHIA COLI.//P21590
 F-NT2RM4002189//MUCIN 2 PRECURSOR (INTESTINAL MUCIN 2).//2.6e-14:233:29//HOMO SAPIENS (HUMAN).//Q02817
 F-NT2RM4002194//TRANSMEMBRANE PROTEIN SEX PRECURSOR.//0.92:108:28//HOMOSAPIENS (HUMAN).//P51805
 F-NT2RM4002205//ELONGATION FACTOR G, MITOCHONDRIAL PRECURSOR (MEF-G).//5.8e-39:122:72//RATTUS NORVEGICUS (RAT).//Q07803
 F-NT2RM4002213//HYPOTHETICAL 88.4 KD PROTEIN B046 4.7 IN CHROMOSOME III.//9.9e-27:110:43//CAENORHABDITIS ELEGANS.//Q03565
 F-NT2RM4002226//GTPASE ACTIVATING PROTEIN ROTUND.//1.3e-21:147:41//DROSOPHILA MELANOGASTER (FRUIT FLY).//P40809
 F-NT2RM4002251//PROTEIN EF-7 (FRAGMENT).//0.00082:45:42//MUS MUSCULUS (MOUSE).//P97805
 F-NT2RM4002256//COLD-REGULATED PROTEIN 1 (FRAGMENT).//0.00015:114:42//HORDEUM VULGARE (BARLEY).//P23251
 F-NT2RM4002266//CUTICLE COLLAGEN 2.//0.00013:142:33//CAENORHABDITIS ELEGANS.//P17656
 F-NT2RM4002278//HYPOTHETICAL 22.2 KD PROTEIN IN NSR1-TIF4631 INTERGENIC REGION.//1.0:40:52//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53288
 F-NT2RM4002281
 F-NT2RM4002287//GAR2 PROTEIN.//0.00055:225:23//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//P41891
 F-NT2RM4002294//HYPOTHETICAL PROTEIN KIAA0281 (HA6725).//1.1e-60:152:75//HOMO SAPIENS (HUMAN).//Q92556
 F-NT2RM4002301//GENERAL STRESS PROTEIN CTC (FRAGMENT).//0.56:43:39//BACILLUS CALDOLYTICUS.//P42832
 F-NT2RM4002323//NONHISTONE CHROMOSOMAL PROTEIN HMG-17.//0.0080:73:35//BOS TAURUS (BOVINE).//P02313
 F-NT2RM4002339//METALLOTHIONEIN 10-III (MT-10-III).//0.67:34:38//MYTILUSEDULIS (BLUE MUSSEL).//P80248
 F-NT2RM4002344//METALLOTHIONEIN-I (MT-I).//0.84:41:31//MUS MUSCULUS (MOUSE).//P02802
 F-NT2RM4002373//GLUTENIN, HIGH MOLECULAR WEIGHT SUBUNIT DY10 PRECURSOR.//0.0019:190:28//TRITICUM AESTIVUM (WHEAT).//P10387
 F-NT2RM4002374//SE5 ANTIGEN.//0.0059:170:32//RATTUS NORVEGICUS (RAT).//Q63003

2986

F-NT2RM4002383//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!/0.13:17:88//HOMO SAPIENS (HUMAN).//P39193
 F-NT2RM4002390
 F-NT2RM4002398//HNRNP ARGININE N-METHYLTRANSFERASE (EC 2.1.1.-) (ODP1 PROTEIN).//0.034:110:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38074
 F-NT2RM4002409//ACETYL-COENZYME A SYNTHETASE (EC 6.2.1.1) (ACETATE--COALIGASE) (ACYL- ACTIVATING ENZYME).//4.0e-20:179:31//METHANOTRIX SOEHNENII.//P27095
 F-NT2RM4002438//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/1.7e-15:41:95//HOMO SAPIENS (HUMAN).//P39194
 F-NT2RM4002446//CRYPTIDIN-RELATED PROTEIN 4C-1 PRECURSOR (CRS4C).//0.0058:24:50//MUS MUSCULUS (MOUSE).//P17534
 F-NT2RM4002452//METALLOTHIONEIN 10-II (MT-10-II).//0.83:48:37//MYTILUS EDULIS (BLUE MUSSEL).//P80247
 F-NT2RM4002457//!!!! ALU SUBFAMILY SC WARNING ENTRY !!!!!/4.9e-07:52:63//HOMO SAPIENS (HUMAN).//P39192
 F-NT2RM4002460//C-HORDEIN (CLONE PC-919) (FRAGMENT).//0.92:43:30//HORDEUM VULGARE (BARLEY).//P17992
 F-NT2RM4002479//RNA HELICASE-LIKE PROTEIN DB10.//1.7e-28:200:41//NICOTIANA SYLVESTRIS (WOOD TOBACCO).//P46942
 F-NT2RM4002482//HYPOTHETICAL 65.9 KD PROTEIN YPRO65W.//8.8e-26:123:49//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q12514
 F-NT2RM4002493//LARVAL CUTICLE PROTEIN I PRECURSOR.//0.17:126:27//DROSOPHILA MIRANDA (FRUIT FLY).//P91627
 F-NT2RM4002499//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!/9.4e-34:92:80//HOMO SAPIENS (HUMAN).//P39194
 F-NT2RM4002504//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!/3.4e-19:55:83//HOMO SAPIENS (HUMAN).//P39189
 F-NT2RM4002527//WD-40 REPEAT PROTEIN MSI2.//3.0e-07:193:27//ARABIDOPSIS THALIANA (MOUSE-EAR CRESS).//022468
 F-NT2RM4002532//AEROLYSIN REGULATORY PROTEIN.//0.97:19:47//AEROMONAS SOBRIA.//P09165
 F-NT2RM4002534//MITOCHONDRIAL 60S RIBOSOMAL PROTEIN L32 PRECURSOR (YML32).//0.76:86:22//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P25348
 F-NT2RM4002558//LONG-CHAIN FATTY ACID TRANSPORT PROTEIN (FATP).//4.2e-55:204:50//MUS MUSCULUS (MOUSE).//Q60714
 F-NT2RM4002565//CHYMOTRYPSIN/ELASTASE ISOINHIBITOR

2987

S 2 TO 5. //1.0:16:62//ASCARIS SUUM (PIG ROUNDWORM) (ASCARIS LUMBRICOIDES). //P07852
 F-NT2RM4002567//HYPOTHETICAL 74.0 KD PROTEIN IN CA J1-HOM3 INTERGENIC REGION. //2.7e-10:184:29//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40032
 F-NT2RM4002571//POLYPEPTIDE N-ACETYL GALACTOSAMINYL TRANSFERASE (EC 2.4.1.41) (PROTEIN- UDP ACETYL GALACTOSAMINYLTRANSFERASE) (UDP-GALNAC: POLYPEPTIDE, N-ACETYL GALACTOSAMINYLTRANSFERASE) (GALNAC-T1). //2.4e-25:124:47//HOMO SAPIENS (HUMAN). //Q10472
 F-NT2RM4002593//HYPOTHETICAL 9.1 KD PROTEIN IN TET B-EXOA INTERGENIC REGION. //0.95:36:38//BACILLUS SUBTILIS. //P37509
 F-NT2RM4002594//MSP1 PROTEIN HOMOLOG. //9.0e-68:227:60//CAENORHABDITIS ELEGANS. //P54815
 F-NT2RM4002623//ASPARTYL-TRNA SYNTHETASE (EC 6.1.1.12) (ASPARTATE--TRNALIGASE) (ASPRS). //3.3e-54:243:47//SYNECHOCYSTIS SP. (STRAIN PCC 6803). //P73851
 F-NT2RP1000018//SUPPRESSOR PROTEIN SRP40. //0.0023:131:25//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32583
 F-NT2RP1000035//RING CANAL PROTEIN (KELCH PROTEIN). //1.0e-06:63:34//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q04652
 F-NT2RP1000040//LETHAL NEUROTOXIN TX1. //0.69:21:47//PHONEUTRIA NIGRIVENTER (BRAZILIAN ARMED SPIDER). //P17727
 F-NT2RP1000063//HYPOTHETICAL 25.1 KD PROTEIN IN SM C3-MRPL8 INTERGENIC REGION. //3.8e-14:130:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40359
 F-NT2RP1000086//HYPOTHETICAL 9.4 KD PROTEIN IN RNP A-THDF INTERGENIC REGION. //0.16:44:40//ESCHERICHIA COLI. //P22847
 F-NT2RP1000101//45.8 KD PROTEIN IN SHM1-MRPL37 INTERGENIC REGION. //1.9e-06:74:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P38344
 F-NT2RP1000111//COP1 REGULATORY PROTEIN (FUSCA PROTEIN FUS1). //2.7e-19:135:36//ARABIDOPSIS THALIANA (MOUSE-EAR CRESS). //P43254
 F-NT2RP1000112//DUAL SPECIFICITY PROTEIN KINASE TT K (EC 2.7.1.-) (PYT). //1.2e-39:91:62//HOMO SAPIENS (HUMAN). //P33981
 F-NT2RP1000124//ATP-DEPENDENT PROTEASE LA 2 (EC 3.4.21.53). //0.074:131:24//MYXOCOCCUS XANTHUS. //P36774
 F-NT2RP1000130//HEPATOMA-DERIVED GROWTH FACTOR (HGF). //1.5e-49:186:56//MUS MUSCULUS (MOUSE). //P51859
 F-NT2RP1000163//METALLOTHIONEIN (MT). //0.98:41:34//PLEURONECTES PLATESSA (PLAICE). //P07216

2988

F-NT2RP1000170//SALIVARY PROLINE-RICH PROTEIN PO (ALLELE K) [CONTAINS: PEPTIDE P-D] (FRAGMENT). //0.85:64:35//HOMO SAPIENS (HUMAN). //P10162
 F-NT2RP1000174//IMMEDIATE-EARLY PROTEIN IE180. //0.00056:89:37//PSEUDORABIES VIRUS (STRAIN KAPLAN) (P RV). //P33479
 F-NT2RP1000191//NIFU PROTEIN. //0.53:78:35//FRANKIA ALNI. //P46045
 F-NT2RP1000202//ANKYRIN, BRAIN VARIANT 2 (ANKYRIN B) (ANKYRIN, NONERYTHROID) (FRAGMENT). //9.1e-21:148:39//HOMO SAPIENS (HUMAN). //Q01485
 F-NT2RP1000243//HYPOTHETICAL PROTEIN MJ1136. //1.4e-37:219:36//METHANOCOCCUS JANNASCHII. //Q58536
 F-NT2RP1000259//HYPOTHETICAL PROTEIN TP0318. //0.18:25:44//TREPONEMA PALLIDUM. //083338
 F-NT2RP1000272//SPLICING FACTOR, ARGININE/SERINE-RICH 3 (PRE-MRNA SPLICING FACTOR SRP20) (X16 PROTEIN). //1.6e-18:133:36//HOMO SAPIENS (HUMAN), AND MUS MUSCULUS (MOUSE). //P23152
 F-NT2RP1000324
 F-NT2RP1000326//HYPOTHETICAL 29.8 KD PROTEIN ZC97.1 IN CHROMOSOME III. //1.0e-23:129:36//CAENORHABDITIS ELEGANS. //P34599
 F-NT2RP1000333//ANTI-SILENCING PROTEIN 1. //2.5e-45:147:57//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32447
 F-NT2RP1000348//REDUCED VIABILITY UPON STARVATION PROTEIN 161. //4.8e-14:119:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P25343
 F-NT2RP1000357//TRYPOMASTIGOTE DECAY-ACCELERATING FACTOR (T-DAF) (FRAGMENT). //1.0:43:32//TRYPANOSOMA CRUZI. //Q26327
 F-NT2RP1000358//HYPOTHETICAL 84.4 KD PROTEIN IN RP C2/RET1 3' REGION. //7.9e-28:244:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P39744
 F-NT2RP1000363//SKIN SECRETORY PROTEIN XP2 PRECURSOR (APEG PROTEIN). //2.2e-07:178:30//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P17437
 F-NT2RP1000376//ANKYRIN R (ANKYRINS 2.1 AND 2.2) (ERYTHROCYTE ANKYRIN). //1.5e-20:254:31//HOMO SAPIENS (HUMAN). //P16157
 F-NT2RP1000409//CYTOCHROME C3 (CYTOCHROME C7) (C55 1.5). //1.0:34:26//DESULFUROMONAS ACETOXIDANS (CHLOROPSEUDOMONAS ETHYLICA). //P00137
 F-NT2RP1000413//MEMBRANE-ASSOCIATED PROTEIN HEM-2 (NAP1 PROTEIN). //3.7e-131:230:97//RATTUS NORVEGICUS (RAT). //P55161
 F-NT2RP1000416//SALIVARY GLUE PROTEIN SCS-3 PRECURSOR. //0.83:54:40//DROSOPHILA SIMULANS (FRUIT FLY). //P13729

2989

F-NT2RP1000418//HYPOTHETICAL 9.9 KD PROTEIN IN GCV
T-SPOIIIAA INTERGENIC REGION. //0.24:91:35//BACILLUS
SUBTILIS. //P49779

F-NT2RP1000439//HYPOTHETICAL 100.5 KD PROTEIN C1B
9.04 IN CHROMOSOME I. //0.13:172:22//SCHIZOSACCHARO
MYCES POMBE (FISSION YEAST). //Q10429

F-NT2RP1000443//QUINONE OXIDOREDUCTASE (EC 1.6.5.
5) (NADPH:QUINONE REDUCTASE) (ZETA- CRYSTALLIN). //
1.9e-08:167:24//HOMO SAPIENS (HUMAN). //Q08257F-NT2
RP1000460//NUCLEAR MOVEMENT PROTEIN NUDC. //1.0e-1
8:149:34//EMERICELLA NIDULANS (ASPERGILLUS NIDULAN
S). //P17624

F-NT2RP1000470//PUTATIVE ATP-DEPENDENT RNA HELICAS
E T26G10.1 IN CHROMOSOME III. //1.3e-43:180:47//CAE
NORHABDITIS ELEGANS. //P34580

F-NT2RP1000478//TUBULIN BETA-6 CHAIN (CLASS-VI). //
1.5e-45:85:63//GALLUSGALLUS (CHICKEN). //P09207

F-NT2RP1000481//HYPOTHETICAL 5.8 KD PROTEIN IN PUH
A 5' REGION (ORF55). //0.083:21:47//RHODOBACTER CAPS
ULATUS (RHODOPSEUDOMONAS CAPSULATA). //P26159F-NT2R
P1000493//POSSIBLE DNA-REPAIR PROTEIN XP-E (POSSIB
LE XERODERMA PIGMENTOSUM GROUP E PROTEIN) (UV-DAMA
GED DNA-BINDING PROTEIN) (UV-DDB). //6.6e-11:139:31
//CERCOPIITHECUS AETHIOPS (GREEN MONKEY) (GRIVET). //
P33194

F-NT2RP1000513//60S RIBOSOMAL PROTEIN L22. //0.017:
92:30//DROSOPHILA MELANOGASTER (FRUIT FLY). //P5088
7

F-NT2RP1000522//UBIQUITIN CARBOXYL-TERMINAL HYDROL
ASE DUB-1 (EC 3.1.2.15) (UBIQUITIN THIOLESTERASE D
UB-1) (UBIQUITIN-SPECIFIC PROCESSING PROTEASE DUB-
1) (DEUBIQUITINATING ENZYME 1). //0.0055:86:36//MUS
MUSCULUS (MOUSE). //Q61068

F-NT2RP1000547//COP-COATED VESICLE MEMBRANE PROTEI
N P24 PRECURSOR (FRAGMENT). //1.2e-09:69:36//CRICET
ULUS GRISEUS (CHINESE HAMSTER). //P49020

F-NT2RP1000574//HOMEBOX PROTEIN MEIS2 (MEIS1-RELA
TED PROTEIN 1). //6.0e-39:141:65//MUS MUSCULUS (MOU
SE). //P97367

F-NT2RP1000577//PUTATIVE ATP-DEPENDENT RNA HELICAS
E YDLO31W. //0.00016:48:45//SACCHAROMYCES CEREVISIA
E (BAKER'S YEAST). //Q12389

F-NT2RP1000581//VON WILLEBRAND FACTOR PRECURSOR. //
0.00017:61:50//HOMO SAPIENS (HUMAN). //P04275

F-NT2RP1000609//LINOLEOYL-COA DESATURASE (EC 1.14.
99.25) (DELTA(6)-DESATURASE). //4.4e-07:128:31//SYN
ECHOCYSTIS SP. (STRAIN PCC 6803). //Q08871

F-NT2RP1000629//CLATHRIN COAT ASSEMBLY PROTEIN AP4
7 (CLATHRIN COAT ASSOCIATED PROTEIN AP47) (GOLGI A
DAPTOR AP-1 47 KD PROTEIN) (HA1 47 KD SUBUNIT) (CL

2990

ATHRIN ASSEMBLY PROTEIN ASSEMBLY PROTEIN COMPLEX 1
MEDIUM CHAIN). //4.2e-70:167:86//MUS MUSCULUS (MOU
SE). //P35585

F-NT2RP1000630//HYPOTHETICAL 118.4 KD PROTEIN IN B
AT2-DAL5 INTERGENIC REGION PRECURSOR. //0.0011:238:
21//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P47
179

F-NT2RP1000677//COLLAGEN ALPHA 1(XVI) CHAIN PRECUR
SOR. //0.99:71:33//HOMOSAPIENS (HUMAN). //Q07092

10 F-NT2RP1000688//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/0.0024:19:94//HOMO SAPIENS (HUMAN). //P3919
3

F-NT2RP1000695//HYPOTHETICAL 83.8 KD PROTEIN C27F
2.7 IN CHROMOSOME III. //2.2e-30:185:37//CAENORHABD
ITIS ELEGANS. //Q18262

F-NT2RP1000701//PHOSPHOLIPASE A-2-ACTIVATING PROTE
IN (PLAP). //3.2e-65:128:93//RATTUS NORVEGICUS (RA
T). //P54319

20 F-NT2RP1000721//TRANSCRIPTION-INITIATION FACTOR TF
IID 135 KD SUBUNIT (TAFII-135) (TAFII135). (TAFII-1
30) (TAFII130). //2.3e-06:139:34//HOMO SAPIENS (HUM
AN). //000268

F-NT2RP1000730//MYOSIN LIGHT CHAIN 1, SLOW-TWITCH
MUSCLE B/VENTRICULAR ISOFORM (FRAGMENT). //0.89:40:
40//MUS MUSCULUS (MOUSE). //P09542

F-NT2RP1000733//METALLOTHIONEIN-LIKE PROTEIN CRS5.
//0.024:24:45//SACCHAROMYCES CEREVISIAE (BAKER'S Y
EAST). //P41902

F-NT2RP1000738//SALIVARY ACIDIC PROLINE-RICH PHOSP
HOPROTEIN 1/2 PRECURSOR (PRP-1 / PRP-3) (PRP-2 / P
RP-4) (PIF-F / PIF-S) (PROTEIN A / PROTEIN C) [CON
TAINS: PEPTIDE P-C]. //0.040:82:36//HOMO SAPIENS (H
UMAN). //P02810

F-NT2RP1000746//HYPOTHETICAL 27.1 KD PROTEIN UFD4-
CAP1 INTERGENIC REGION. //2.0e-30:170:37//SACCHAROM
YCES CEREVISIAE (BAKER'S YEAST). //P33201

F-NT2RP1000767//PSEUDOMONAPEPSIN PRECURSOR (EC 3.
4.23.37) (PEPSTATIN-INSENSITIVE CARBOXYL PROTEINASE
E). //0.99:75:34//PSEUDOMONAS SP. (STRAIN 101). //P4
2790

F-NT2RP1000782//CELL SURFACE GLYCOPROTEIN A15 (T-C
ELL ACUTE LYMPHOBLASTIC LEUKEMIA ASSOCIATED ANTIGE
N 1) (TALLA-1) (MEMBRANE COMPONENT, X CHROMOSOME,
SURFACE MARKER 1). //2.3e-23:159:35//HOMO SAPIENS
(HUMAN). //P41732

F-NT2RP1000796//CORNIFIN (SMALL PROLINE-RICH PROTE
IN I) (SPR-I) (SMALL PROLINE-RICH SQUAMOUS CELL MA
RKER) (SPRP). //0.00018:79:32//SUS SCROFA (PIG). //P
35323

F-NT2RP1000825//GTPASE-ACTIVATING PROTEIN RHOGAP

2991

(RHO-RELATED SMALL GTPASE PROTEIN ACTIVATOR) (CDC4 2 GTPASE-ACTIVATING PROTEIN) (P50-RHOGAP). //3.1e-3 7:89:64//HOMO SAPIENS (HUMAN). //Q07960

F-NT2RP1000833//SMALL PROLINE RICH PROTEIN II (SPR-II) (CLONE 930). //0.32:29:48//HOMO SAPIENS (HUMAN). //P22531

F-NT2RP1000834//2-ARYLPROPIONYL-COA EPIMERASE (EC 5.-.-.-). //6.4e-67:202:68//RATTUS NORVEGICUS (RAT). //P70473

F-NT2RP1000836//HYPOTHETICAL 7.3 KD PROTEIN IN 100 KD PROTEIN REGION. //1.0:35:54//HUMAN ADENOVIRUS TYPE 41. //P23691

F-NT2RP1000846//SMALL PROLINE-RICH PROTEIN 2-1. //0.013:35:48//HOMO SAPIENS (HUMAN). //P35326

F-NT2RP1000851//PERIOD CLOCK PROTEIN (FRAGMENT). //0.082:28:57//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q04536

F-NT2RP1000856//PLATELET-ENDOTHELIAL TETRASPAN ANTIGEN 3 (PETA-3) (GP27) (MEMBRANE GLYCOPROTEIN SFA-1) (CD151 ANTIGEN). //2.5e-26:190:30//MUS MUSCULUS (MOUSE). //O35566

F-NT2RP1000860//POTENTIAL TRANSCRIPTIONAL ADAPTOR. //0.13:86:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q02336

F-NT2RP1000902//HYPOTHETICAL 127.4 KD PROTEIN F07F 6.4 IN CHROMOSOME III. //7.6e-11:200:35//CAENORHABDITIS ELEGANS. //Q09531

F-NT2RP1000915//HYPOTHETICAL GTP-BINDING PROTEIN IN PMI40-PAC2 INTERGENIC REGION. //1.4e-06:88:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40010

F-NT2RP1000916//SUPPRESSOR PROTEIN SRP40. //0.40:90:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P32583

F-NT2RP1000943//MUCIN 2 PRECURSOR (INTESTINAL MUCIN 2). //0.099:75:34//HOMO SAPIENS (HUMAN). //Q02817

F-NT2RP1000944//HYPOTHETICAL 29.3 KD PROTEIN (ORF9 2). //7.6e-06:65:41//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPMNPV). //O10341

F-NT2RP1000947//UBIQUITIN-CONJUGATING ENZYME E2-17 KD 2 (EC 6.3.2.19) (UBIQUITIN-PROTEIN LIGASE) (UBIQUITIN CARRIER PROTEIN) (E2(17)KB 2). //3.6e-12:27:77//HOMO SAPIENS (HUMAN), MUS MUSCULUS (MOUSE), RATTUS NORVEGICUS (RAT), AND XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P51669

F-NT2RP1000954//RING CANAL PROTEIN (KELCH PROTEIN). //2.8e-15:169:28//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q04652

F-NT2RP1000958//HYPOTHETICAL GTP-BINDING PROTEIN IN PMI40-PAC2 INTERGENIC REGION. //4.2e-16:162:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40010

2992

F-NT2RP1000959//CORNIFIN A (SMALL PROLINE-RICH PROTEIN IA) (SPR-IA) (SPRK). //0.0031:34:44//HOMO SAPIENS (HUMAN). //P35321

F-NT2RP1000966//NUCLEOLIN (PROTEIN C23). //1.5e-52:110:95//HOMO SAPIENS (HUMAN). //P19338

F-NT2RP1000980//LIGHT-HARVESTING PROTEIN B-1015, A LPHA CHAIN PRECURSOR (ANTENNA PIGMENT PROTEIN, ALPHA CHAIN). //0.87:37:45//RHODOSPIRILLUM RUBRUM (RED SPHERULE). //P04123

F-NT2RP1000988

F-NT2RP1001011//PROTEIN P19. //0.96:30:50//BACTERIOPHAGE PRD1. //P17638

F-NT2RP1001013//DNA-BINDING PROTEIN 65 (PROTEIN GP 65). //1.0:20:45//BACTERIOPHAGE T4. //P16012

F-NT2RP1001014

F-NT2RP1001033//TUBULIN GAMMA CHAIN. //2.5e-16:112:42//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //P25295

F-NT2RP1001073//HYPOTHETICAL 10.4 KD PROTEIN IN FT R1-SPT15 INTERGENIC REGION. //7.6e-16:82:51//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40089

F-NT2RP1001079//SARCOSINE OXIDASE (EC 1.5.3.1). //4.8e-15:95:40//ARTHROBACTER SP. (STRAIN TE1826). //P40873

F-NT2RP1001080//PROBABLE ATP-DEPENDENT RNA HELICASE DBP9. //2.4e-29:126:46//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q06218

F-NT2RP1001113//SMALL PROLINE-RICH PROTEIN 2-1. //0.49:38:39//HOMO SAPIENS (HUMAN). //P35326

30 F-NT2RP1001173//RHOMBOTIN-1 (CYSTEINE RICH PROTEIN TTG-1) (T-CELL TRANSLOCATION PROTEIN 1) (LIM-ONLY PROTEIN 1). //0.99:54:37//HOMO SAPIENS (HUMAN). //P25800

F-NT2RP1001177//HISTONE MACRO-H2A.1. //1.6e-29:85:76//RATTUS NORVEGICUS (RAT). //Q02874

F-NT2RP1001185

F-NT2RP1001199//NEUROTOXIN I. //1.0:23:47//CENTRUROIDES SCULPTURATUS (BARK SCORPION). //P01491

F-NT2RP1001247//TRANSFORMING GROWTH FACTOR BETA 4 PRECURSOR (TGF-BETA 4) (ENDOMETRIAL BLEEDING-ASSOCIATED FACTOR). //3.3e-08:28:89//HOMO SAPIENS (HUMAN). //O00292

F-NT2RP1001248//TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN). //0.33:49:28//HUMAN IMMUNODEFICIENCY VIRUS TYPE 1 (NDK ISOLATE) (HIV-1). //P18804

F-NT2RP1001253//GLUCOSAMINE-6-PHOSPHATE ISOMERASE (EC 5.3.1.10) (GLUCOSAMINE-6-PHOSPHATE DEAMINASE) (GNPDA) (OSCILLIN) (KIAA0060). //3.8e-46:115:81//HOMO SAPIENS (HUMAN). //P46926

50 F-NT2RP1001286//GALECTIN-3 (GALACTOSE-SPECIFIC LEC

- TIN 3) (MAC-2 ANTIGEN) (ICE-BINDING PROTEIN) (35 KD LECTIN) (CARBOHYDRATE BINDING PROTEIN 35) (CBP 35) (LAMININ-BINDING PROTEIN) (LECTIN L-29) (L-34 GAL ACTOSIDE-BINDING LECTIN). //0.16:48:37//MUS MUSCULUS (MOUSE). //P16110
- F-NT2RP1001294//MICROTUBULE-ASSOCIATED PROTEIN YTM 1. //6.1e-05:92:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q12024
- F-NT2RP1001302//MICROTUBULE-ASSOCIATED PROTEIN YTM 1. //1.2e-05:92:33//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q12024
- F-NT2RP1001310//PROBABLE E4 PROTEIN. //0.99:109:26//HUMAN PAPILLOMAVIRUS TYPE 5. //P06924
- F-NT2RP1001311//SODIUM/HYDROGEN EXCHANGER 5 (NA(+)/H(+) EXCHANGER 5) (NHE-5) (FRAGMENT). //0.99:94:31//HOMO SAPIENS (HUMAN). //Q14940
- F-NT2RP1001313//CYTOCHROME B5. //9.0e-13:92:38//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40312
- F-NT2RP1001361//NADH-UBIQUINONE OXIDOREDUCTASE SUBUNIT B14.5B (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-B14.5B) (CI-B14.5B). //1.2e-47:117:74//BOS TAURUS (BOVINE). //Q02827
- F-NT2RP1001385//CELL DIVISION PROTEIN FTSN. //0.64:107:28//ESCHERICHIA COLI. //P29131
- F-NT2RP1001395//PROCOLLAGEN ALPHA 2(I) CHAIN PRECURSOR (FRAGMENTS). //0.25:35:45//GALLUS GALLUS (CHICKEN). //P02467
- F-NT2RP1001410//PUTATIVE GTP-BINDING PROTEIN W08E 3.3. //2.2e-41:129:67//CAENORHABDITIS ELEGANS. //P91917
- F-NT2RP1001424//UREASE ACCESSORY PROTEIN UREF (FRAGMENT). //0.87:24:45//ESCHERICHIA COLI. //Q03286
- F-NT2RP1001432//CYSTEINE PROTEINASE INHIBITOR B (CYSTATIN B) (SCB). //1.0:35:42//HELIANTHUS ANNUUS (COMMON SUNFLOWER). //Q10993
- F-NT2RP1001449//KERATIN, ULTRA HIGH-SULFUR MATRIX PROTEIN (UHS KERATIN). //0.053:37:37//OVIS ARIES (SHEEP). //P26372
- F-NT2RP1001457//HYPOTHETICAL 57.0 KD TRP-ASP REPEATS CONTAINING PROTEININ CPR4-SSK22 INTERGENIC REGION. //2.9e-16:159:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P25382
- F-NT2RP1001466//HYPOTHETICAL PROTEIN MJ0284. //5.3e-15:162:35//METHANOCOCCUS JANNASCHII. //Q57732
- F-NT2RP1001475//HYPOTHETICAL 195.1 KD PROTEIN IN DNA43-UBI1 INTERGENIC REGION. //0.69:119:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40457
- F-NT2RP1001482//PROTEASOME COMPONENT C9 (EC 3.4.9.46) (MACROPAIN SUBUNIT C9) (MULTICATALYTIC ENDOPEPTIDASE COMPLEX SUBUNIT C9). //1.0:58:32//HOMO SAPIENS (HUMAN). //P25789
- F-NT2RP1001494//MALE STERILITY PROTEIN 2. //2.4e-12:84:42//ARABIDOPSIS THALIANA (MOUSE-EAR CRESS). //Q08891
- F-NT2RP1001543//MYO-INOSITOL-1-PHOSPHATE SYNTHASE (EC 5.5.1.4) (IPS). //6.3e-37:94:52//SPIRODELA POLYRRHIZA. //P42803
- F-NT2RP1001546//LEUKOCYTE SURFACE ANTIGEN CD53 (CELL SURFACE GLYCOPROTEIN CD53). //9.3e-11:98:29//HOMO SAPIENS (HUMAN). //P19397
- F-NT2RP1001569//SIGNAL RECOGNITION PARTICLE RECEPTOR BETA SUBUNIT (SR-BETA). //2.2e-64:159:84//MUS MUSCULUS (MOUSE). //P47758
- F-NT2RP1001616//HYPOTHETICAL 13.5 KD PROTEIN C45G 9.7 IN CHROMOSOME III. //9.2e-05:49:42//CAENORHABDITIS ELEGANS. //Q09506
- F-NT2RP1001665//REGB PROTEIN. //0.99:29:37//PSEUDOMONAS AERUGINOSA. //Q03381
- F-NT2RP2000001//SMALL PROLINE-RICH PROTEIN 2-1. //0.64:36:41//HOMO SAPIENS (HUMAN). //P35326
- F-NT2RP2000006//DNAJ PROTEIN HOMOLOG 1 (HDJ-1) (HEAT SHOCK PROTEIN 40) (HSP40). //1.7e-19:74:52//HOMO SAPIENS (HUMAN). //P25685
- F-NT2RP2000007//TROPOMYOSIN, FIBROBLAST AND EPITHELIAL MUSCLE-TYPE (TM36) (TME1) (TM1). //0.93:126:23//HOMO SAPIENS (HUMAN). //P06468
- F-NT2RP2000008//ZINC FINGER PROTEIN 33A (ZINC FINGER PROTEIN KOX31) (KIAA0065) (HA0946) (FRAGMENT). //4.2e-35:156:54//HOMO SAPIENS (HUMAN). //Q06730
- F-NT2RP2000027//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 5 (EC 1.6.5.3) (FRAGMENT). //0.95:41:39//MACACA FASCICULARIS (CRAB EATING MACAQUE) (CYNOMOLGUS MONKEY). //P50665
- F-NT2RP2000032//BAX PROTEIN, CYTOPLASMIC ISOFORM GAMMA. //1.0:35:34//HOMO SAPIENS (HUMAN). //Q07815
- F-NT2RP2000040//BASIC PROLINE-RICH PEPTIDE IB-1. //0.0024:58:36//HOMO SAPIENS (HUMAN). //P04281
- F-NT2RP2000045//DNAJ PROTEIN. //1.1e-12:42:66//THERMOPHILUS AQUATICUS (SUBSP. THERMOPHILUS). //Q56237
- F-NT2RP2000054//GONADOLIBERIN III PRECURSOR (GONADOTROPIN-RELEASING HORMONE III) (GNRH-III) (LH-RH III) (LULIBERIN III). //0.20:46:36//ONCORHYNCHUS MASOU (CHERRY SALMON) (MASU SALMON). //P30973
- F-NT2RP2000056//PROTEIN-TYROSINE PHOSPHATASE EPSILON PRECURSOR (EC 3.1.3.48) (R-PTP- EPSILON). //1.3e-18:45:100//MUS MUSCULUS (MOUSE). //P49446
- F-NT2RP2000067//HOMEODOMAIN PROTEIN HOX-A5 (S12-B) (FRAGMENT). //0.71:44:40//SALMO SALAR (ATLANTIC SALMON). //P09637
- F-NT2RP2000070//INSULIN. //0.94:30:43//HYSTRIX CRISTATUS (HUMAN). //P25789

2995

TATA (CRESTED PORCUPINE). //P01328
 F-NT2RP2000076//ETS-LIKE PROTEIN POINTED P1 (D-ETS-2). //0.0013:76:40//DROSOPHILA MELANOGASTER (FRUIT FLY). //P51022
 F-NT2RP2000077//U1 SMALL NUCLEAR RIBONUCLEOPROTEIN C (U1-C). //0.24:49:40//HOMO SAPIENS (HUMAN). //P09234
 F-NT2RP2000079//PLATELET FACTOR 4 (PF-4). //0.15:52:30//SUS SCROFA (PIG). //P30034
 F-NT2RP2000088//HYPOTHETICAL 13.6 KD PROTEIN IN SP T4-ROM1 INTERGENIC REGION. //1.0:36:44//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53245
 F-NT2RP2000091//HYPOTHETICAL PROTEIN HI0149 PRECURSOR. //0.22:38:47//HAEMOPHILUS INFLUENZAE. //P43953
 F-NT2RP2000097//VIRUS ATTACHMENT PROTEIN (O61R). //0.75:33:36//AFRICAN SWINE FEVER VIRUS (STRAIN BA71 V) (ASFV). //P32510
 F-NT2RP2000098
 F-NT2RP2000108//IIII ALU SUBFAMILY SX WARNING ENTRY IIII//2.4e-09:50:70//HOMO SAPIENS (HUMAN). //P39195
 F-NT2RP2000114//WISKOTT-ALDRICH SYNDROME PROTEIN (WASP). //0.024:52:44//HOMO SAPIENS (HUMAN). //P42768
 F-NT2RP2000120//5.8 KD PROTEIN IN HMC OPERON (ORF 4). //0.67:37:32//DESULFOVIBRIO VULGARIS (STRAIN HILDENBOROUGH). //P33391
 F-NT2RP2000126//CHROMODOMAIN-HELICASE-DNA-BINDING PROTEIN 1 (CHD-1). //1.5e-23:94:47//HOMO SAPIENS (HUMAN). //014646
 F-NT2RP2000133//SPLICEOSOME ASSOCIATED PROTEIN 49 (SAP 49) (SF3B53). //5.6e-10:82:39//HOMO SAPIENS (HUMAN). //Q15427
 F-NT2RP2000147//CLATHRIN COAT ASSEMBLY PROTEIN AP47 (CLATHRIN COAT ASSOCIATED PROTEIN AP47) (COLGI ADAPTOR AP-1 47 KD PROTEIN) (HA1 47 KD SUBUNIT) (CLATHRIN ASSEMBLY PROTEIN ASSEMBLY PROTEIN COMPLEX 1 MEDIUM CHAIN). //6.7e-89:96:98//MUS MUSCULUS (MOUSE). //P35585
 F-NT2RP2000153//PEPTIDYLPROLYL ISOMERASE CYP-1 (EC 5.2.1.8) (PEPTIDYLPROLYL CIS-TRANS ISOMERASE) (CYCLOPHILIN) (PPIASE). //1.7e-05:136:33//BRUGIAMALAYI. //Q27450
 F-NT2RP2000157//ML02 PROTEIN. //2.7e-06:62:40//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09329
 F-NT2RP2000161//DIS3 PROTEIN HOMOLOG. //2.7e-33:173:45//CAENORHABDITIS ELEGANS. //Q17632
 F-NT2RP2000173//HYPOTHETICAL 10.5 KD PROTEIN IN SODA-COMGA INTERGENIC REGION. //0.99:62:25//BACILLUS SUBTILIS. //P54499

2996

F-NT2RP2000175//MALE SPECIFIC SPERM PROTEIN MST84D B. //0.19:41:43//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q01643
 F-NT2RP2000183//DIHYDROPYRIMIDINASE RELATED PROTEIN N-2 (DRP-2) (NEURAL SPECIFIC PROTEIN NSP60). //4.1e-19:114:44//BOS TAURUS (BOVINE). //002675
 F-NT2RP2000195//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (AGL). //0.99:30:33//MICROTUS PENNSYLVANICUS (MEADOW VOLE). //P24949
 F-NT2RP2000205//MERCURIC TRANSPORT PROTEIN PERIPLASMIC COMPONENT PRECURSOR (PERIPLASMIC MERCURY ION BINDING PROTEIN) (MERCURY SCAVENGER PROTEIN). //0.098:88:25//SHEWANELLA PUTREFACIENS (PSEUDOMONAS PUTREFACIENS). //Q54463
 F-NT2RP2000208//MALE SPECIFIC SPERM PROTEIN MST84D D. //0.020:19:57//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q01645
 F-NT2RP2000224//PUTATIVE CUTICLE COLLAGEN C09G5.4. //0.0058:159:32//CAENORHABDITIS ELEGANS. //Q09455
 F-NT2RP2000232//P55-C-FOS PROTO-ONCOGENE PROTEIN (FRAGMENT). //1.0:44:38//OVIS ARIES (SHEEP). //002761
 F-NT2RP2000233//GASTRIN/CHOLECYSTOKININ TYPE B RECEPTOR (CCK-B RECEPTOR) (CCK-BR). //0.34:53:43//CANIS FAMILIARIS (DOG). //P30552
 F-NT2RP2000239//ACIDIC PROLINE-RICH PROTEIN PRECURSOR (CLONE PRP25) (FRAGMENT). //0.019:69:33//RATTUS NORVEGICUS (RAT). //P10164
 F-NT2RP2000248//OVOMUCOID (FRAGMENT). //0.88:18:55//POLYPLECTRON EMPHANUM (PALAWAN PEACOCK-PHEASANT). //P52250
 F-NT2RP2000257//PUTATIVE MITOCHONDRIAL CARRIER YIL006W. //6.4e-09:83:37//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40556
 F-NT2RP2000258//MYOSIN II HEAVY CHAIN, NON MUSCLE. //0.081:217:28//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD). //P08799
 F-NT2RP2000270//IIII ALU SUBFAMILY J WARNING ENTRY IIII//2.4e-17:80:57//HOMO SAPIENS (HUMAN). //P39188
 F-NT2RP2000274//HYPOTHETICAL 5.8 KD PROTEIN. //0.082:22:45//CLOVER YELLOW MOSAIC VIRUS (CYMV). //P16485
 F-NT2RP2000283//HYPOTHETICAL 83.6 KD PROTEIN R05D 3.2 IN CHROMOSOME III. //0.39:38:34//CAENORHABDITIS ELEGANS. //P34535
 F-NT2RP2000288
 F-NT2RP2000289//HYPOTHETICAL 9.4 KD PROTEIN IN RNP A-THDF INTERGENIC REGION. //0.40:38:42//ESCHERICHIA COLI. //P22847
 F-NT2RP2000297//ZINC FINGER PROTEIN 85 (ZINC FINGER

2997

R PROTEIN HPF4) (HTF1). //2.3e-62:206:47//HOMO SAPIENS (HUMAN). //Q03923
 F-NT2RP2000298//CUTICLE COLLAGEN 12 PRECURSOR. //0.55:81:40//CAENORHABDITIS ELEGANS. //P20630
 F-NT2RP2000310//RUBREDOXIN (RD). //0.13:43:41//TREPONEMA PALLIDUM. //083956
 F-NT2RP2000327//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (A6L). //1.0:46:30//CADUS MORHUA (ATLANTIC COD). //P15996
 F-NT2RP2000328//HYPOTHETICAL 86.6 KD PROTEIN IN PF 10 K1-TDS4 INTERGENIC REGION. //2.0e-21:198:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53313
 F-NT2RP2000329//GTP:AMP PHOSPHOTRANSFERASE MITOCHONDRIAL (EC 2.7.4.10) (AK3). //1.8e-91:155:92//BOS TAURUS (BOVINE). //P08760
 F-NT2RP2000337//PROTEIN A54. //0.75:48:35//VACCINIA VIRUS (STRAIN WR), AND VACCINIA VIRUS (STRAIN COPENHAGEN). //P21072
 F-NT2RP2000346//MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD116. //9.7e-13:114:42//MUS MUSCULUS (MOUSE). //P17564
 F-NT2RP2000369//CALTRIN (CALCIUM TRANSPORT INHIBITOR). //0.98:47:34//MUS MUSCULUS (MOUSE). //Q09098
 F-NT2RP2000412//SHORT NEUROTOXIN D PRECURSOR. //0.66:57:36//AIPYSURUS LAEVIS (OLIVE SEA SNAKE). //P19960
 F-NT2RP2000414//HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN F (HNRNP F). //1.0e-27:96:67//HOMO SAPIENS (HUMAN). //P52597
 F-NT2RP2000420//ZINC FINGER PROTEIN 191. //0.16:47:38//HOMO SAPIENS (HUMAN). //014754
 F-NT2RP2000422//PUTATIVE PHOSPHOACETYLGLUCOSAMINE MUTASE (EC 5.4.2.3) (ACETYLGLUCOSAMINE PHOSPHOMUTASE) (N-ACETYLGLUCOSAMINE-PHOSPHATE MUTASE). //3.6e-19:148:36//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09687
 F-NT2RP2000438//TUBULIN GAMMA CHAIN. //0.86:190:27//RETICULOMYXA FILOSA. //P54405
 F-NT2RP2000448//OXYSTEROL-BINDING PROTEIN. //3.7e-13:140:42//HOMO SAPIENS (HUMAN). //P22059
 F-NT2RP2000459//NEURONAL PROTEIN 3.1 (P311 PROTEIN). //1.0:45:35//HOMO SAPIENS (HUMAN). //Q16612
 F-NT2RP2000498//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!! //0.062:25:68//HOMO SAPIENS (HUMAN). //P39194
 F-NT2RP2000503
 F-NT2RP2000510//TOXIN IV-5. //1.0:51:33//TITYUS BAHIENSIS (BRAZILIAN SCORPION). //P56608
 F-NT2RP2000516//SLYX PROTEIN. //1.0:52:32//ESCHERICHIA COLI. //P30857
 F-NT2RP2000523//PHORBOLIN I (FRAGMENTS). //1.4e-06: 50

2998

36:47//HOMO SAPIENS (HUMAN). //P31941
 F-NT2RP2000603//ALPHA/BETA-GLIADIN PRECURSOR (PROLAMIN) (CLASS A-III). //0.93:119:26//TRITICUM AESTIVUM (WHEAT). //P04723
 F-NT2RP2000617//SPERM PROTAMINE P1 (CYSTEINE-RICH PROTAMINE). //0.056:16:62//OVIS ARIES (SHEEP), AND CAPRA HIRCUS (GOAT). //P04102
 F-NT2RP2000634//NEDD-4 PROTEIN (EC 6.3.2.-) (KIAA0093) (FRAGMENT). //1.8e-05:128:28//HOMO SAPIENS (HUMAN). //P46934
 F-NT2RP2000644//HYPOTHETICAL PROTEIN HI1566 PRECURSOR. //0.85:48:39//HAEMOPHILUS INFLUENZAE. //P44257
 F-NT2RP2000656//EARLY GROWTH RESPONSE PROTEIN 1 (EGR-1) (NERVE GROWTH FACTOR-INDUCED PROTEIN A) (NGF I-A). //1.0:111:24//RATTUS NORVEGICUS (RAT). //P08154
 F-NT2RP2000658//URONATE ISOMERASE (EC 5.3.1.12) (GLUCURONATE ISOMERASE) (URONIC ISOMERASE). //0.49:79:31//ESCHERICHIA COLI. //P42607
 F-NT2RP2000668//MEROZOITE SURFACE ANTIGEN 2 PRECURSOR (MSA-2) (45 KD MEROZOITE SURFACE ANTIGEN). //0.020:115:30//PLASMODIUM FALCIPARUM (ISOLATE 3D7). //P50498
 F-NT2RP2000678//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!! //0.00085:38:68//HOMO SAPIENS (HUMAN). //P39188
 F-NT2RP2000704//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!! //2.2e-17:55:74//HOMO SAPIENS (HUMAN). //P39188
 F-NT2RP2000710//ASPARTYL-TRNA SYNTHETASE (EC 6.1.1.12) (ASPARTATE--TRNALIGASE) (ASPRS). //8.9e-47:106:59//TREPONEMA PALLIDUM. //083950
 F-NT2RP2000715
 F-NT2RP2000731//CONIDIATION-SPECIFIC PROTEIN 10. //0.094:31:41//NEUROSPORA CRASSA. //P10713
 F-NT2RP2000758//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!! //0.00027:31:74//HOMO SAPIENS (HUMAN). //P39188
 F-NT2RP2000764//NIFS PROTEIN. //2.7e-27:175:47//ANABAENA SP. (STRAIN PCC7120). //P12623
 F-NT2RP2000809//HYPOTHETICAL PROTEIN MG381 HOMOLOG. //0.91:85:25//MYCOPLASMA PNEUMONIAE. //P75219
 F-NT2RP2000812//DILUTE MYOSIN HEAVY CHAIN, NON-MUSCLE (MYOSIN 5A). //2.8e-07:133:31//MUS MUSCULUS (MOUSE). //Q99104
 F-NT2RP2000814//40S RIBOSOMAL PROTEIN S27A. //0.93:44:38//LYCOPERSICON ESCULENTUM (TOMATO), AND SOLANUM TUBEROSUM (POTATO). //P27083
 F-NT2RP2000816//HYPOTHETICAL 88.4 KD PROTEIN B046 4.7 IN CHROMOSOME III. //3.3e-21:123:39//CAENORHABD

- ITIS ELEGANS.//Q03565
 F-NT2RP2000819//TROPOMYOSIN 5, CYTOSKELETAL TYPE.//
 /1.0:71:30//MUS MUSCULUS (MOUSE).//P21107
 F-NT2RP2000841//GUANINE NUCLEOTIDE RELEASING PROTEIN (GNRP).//0.0011:133:26//MUS MUSCULUS (MOUSE).//P27671
 F-NT2RP2000842//LYSOPHOSPHATIDIC ACID RECEPTOR (EDG-2).//6.4e-13:22:95//HOMO SAPIENS (HUMAN).//Q92633
 F-NT2RP2000845//BOWMAN-BIRK TYPE PROTEINASE INHIBITOR (MSTI).//0.92:24:41//MEDICAGO SCUTELLATA (SNAIL MEDIC).//P80321
 F-NT2RP2000863//N-MYC PROTO-ONCOGENE PROTEIN.//0.010:148:27//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P24793
 F-NT2RP2000880//PROBABLE TRANSLATION INITIATION FACTOR IF-2.//4.0e-100:199:94//HOMO SAPIENS (HUMAN).//060841
 F-NT2RP2000892//PROCOLLAGEN ALPHA 1(II) CHAIN PRECURSOR [CONTAINS: CHONDROCALCIN].//0.43:45:44//MUS MUSCULUS (MOUSE).//P28481
 F-NT2RP2000931//MATRIN 3.//2.8e-46:104:92//RATTUS NORVEGICUS (RAT).//P43244
 F-NT2RP2000932//2-5A-DEPENDENT RIBONUCLEASE (EC 3.1.26.-). (2-5A-DEPENDENT RNAASE) (RNASE L) (RIBONUCLEASE 4) (FRAGMENT).//3.9e-07:113:31//MUS MUSCULUS (MOUSE).//Q05921
 F-NT2RP2000938//VOLTAGE-GATED POTASSIUM CHANNEL PROTEIN KV3.3 (KSH11D).//0.026:59:45//RATTUS NORVEGICUS (RAT).//Q01956
 F-NT2RP2000943//HYPOTHETICAL PROTEIN KIAA0079 (HA3543).//5.9e-18:161:42//HOMO SAPIENS (HUMAN).//P53992
 F-NT2RP2000965//INNER CENTROMERE PROTEIN (INCENP).//0.062:156:25//GALLUSGALLUS (CHICKEN).//P53352
 F-NT2RP2000970//EC PROTEIN HOMOLOG.//1.0:50:30//ARABIDOPSIS THALIANA (MOUSE-EAR CRESS).//P93746
 F-NT2RP2000985//HYPOTHETICAL 96.8 KD PROTEIN IN SI-S2-MTD1 INTERGENIC REGION.//2.5e-06:53:47//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P36159
 F-NT2RP2000987//INSECT TOXIN 4 (INSECT TOXIN AAH I T4).//1.0:32:34//ANDROCTONUS AUSTRALIS HECTOR (SAHARA SCORPION).//P21150
 F-NT2RP2001036//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//1.2e-33:65:81//HOMO SAPIENS (HUMAN).//P39193
 F-NT2RP2001044//HIRUSTASIN.//0.97:15:66//HIRUDO MEDICINALIS (MEDICINAL LEECH).//P80302
 F-NT2RP2001056//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//1.0e-24:85:65//HOMO SAPIENS (HUMAN).//P39194
 F-NT2RP2001065//BOWMAN-BIRK TYPE SEED TRYPSIN AND CHYMOTRYPSIN INHIBITOR(BTCI).//0.41:50:32//VIGNA UNGUICULATA (COWPEA).//P17734
 F-NT2RP2001070//PROBABLE PYRIDOXAMINE 5'-PHOSPHATE OXIDASE (EC 1.4.3.5) (PNP/PMP OXIDASE) (FPRA PROTEIN).//6.2e-18:64:48//MYXOCOCCUS XANTHUS.//P21159
 F-NT2RP2001081//SYNAPTOTAGMIN IV.//7.8e-16:94:46//RATTUS NORVEGICUS (RAT).//P50232
 F-NT2RP2001094//METALLOTHIONEIN-I (MT-I).//1.0:24:33//RATTUS NORVEGICUS(RAT).//P02803
 F-NT2RP2001119//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//7.5e-11:61:63//HOMO SAPIENS (HUMAN).//P39195
 F-NT2RP2001127//XE169 PROTEIN (SMCX PROTEIN) (FRAGMENTS).//1.0e-47:155:58//MUS MUSCULUS (MOUSE).//P41230
 F-NT2RP2001137//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//0.10:68:39//BOS TAURUS (BOVINE).//P25508
 F-NT2RP2001149//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.1e-13:81:59//HOMO SAPIENS (HUMAN).//P39198
 F-NT2RP2001168//PROTEIN KINASE C SUBSTRATE 80 KD PROTEIN (FRAGMENTS).//0.0071:77:33//RATTUS NORVEGICUS (RAT).//P20468
 F-NT2RP2001173//CYTOSKELETON-ASSOCIATED PROTEIN CKAPI (TUBULIN FOLDING COFACTOR B).//1.0:36:41//HOMO SAPIENS (HUMAN).//Q99426
 F-NT2RP2001174//ZINC FINGER PROTEIN 137.//7.2e-11:65:43//HOMO SAPIENS (HUMAN).//P52743
 F-NT2RP2001196//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 2 (EC 1.6.5.3).//1.0:95:26//CAPRA HIRCUS (GOAT).//Q36346
 F-NT2RP2001218//HYPOTHETICAL 59.2 KD PROTEIN IN M0B1-SGA1 INTERGENIC REGION.//0.00024:80:23//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P40486F-NT2RP2001226//RABPHILIN-3A (FRAGMENT).//4.6e-05:121:39//MUS MUSCULUS (MOUSE).//P47708
 F-NT2RP2001233//ZINC FINGER PROTEIN ZFP-36 (FRAGMENT).//3.2e-61:153:56//HOMO SAPIENS (HUMAN).//P16415
 F-NT2RP2001245//SYNAPTONEMAL COMPLEX PROTEIN 1 (SCP-1 PROTEIN).//4.9e-05:230:21//HOMO SAPIENS (HUMAN).//Q15431
 F-NT2RP2001268//HOMEBOX PROTEIN CEH-32.//0.23:15:9:25//CAENORHABDITIS ELEGANS.//Q23175
 F-NT2RP2001277
 F-NT2RP2001290//BETA-SOLUBLE NSF ATTACHMENT PROTEIN (SNAP-BETA) (SNAP-ALPHA HOMOLOG) (BRAIN PROTEIN I47) (FRAGMENT).//1.0e-86:131:97//MUS MUSCULUS (MO

- USE). //P28663
 F-NT2RP2001295
 F-NT2RP2001312//N-ACETYLGLUCOSAMINE-6-SULFATASE PR
 ECURSOR (EC 3.1.6.14) (G6S) (GLUCOSAMINE-6-SULFATAS
 E). //0.64:80:33//CAPRA HIRCUS (GOAT). //P50426
 F-NT2RP2001327//TUMOR NECROSIS FACTOR, ALPHA-INDUC
 ED PROTEIN 1, ENDOTHELIAL (B12 PROTEIN). //1.0e-36:
 118:65//HOMO SAPIENS (HUMAN). //Q13829
 F-NT2RP2001328//PROBABLE E5 PROTEIN. //1.0:46:41//H
 UMAN PAPILOMAVIRUS TYPE 33. //P06426
 F-NT2RP2001347//!!!! ALU SUBFAMILY SP WARNING ENTR
 Y !!!!!/4.5e-19:66:62//HOMO SAPIENS (HUMAN). //P391
 93
 F-NT2RP2001366//SPERM-SPECIFIC PROTEIN PHI-1. //0.6
 6:55:32//MYTILUS EDULIS (BLUE MUSSEL). //Q04621
 F-NT2RP2001378//VOLTAGE-GATED POTASSIUM CHANNEL PR
 OTEIN KV3.3 (KSHIID) (FRAGMENT). //0.060:78:33//HOM
 O SAPIENS (HUMAN). //Q14003
 F-NT2RP2001381//26S PROTEASE REGULATORY SUBUNIT 8
 (SUG1 HOMOLOG) (XSUG1). //1.0:167:26//XENOPUS LAEVI 20
 S (AFRICAN CLAWED FROG). //P46470
 F-NT2RP2001392//KERATIN, HIGH-SULFUR MATRIX PROTEI
 N, IIIA3. //0.0080:82:32//OVIS ARIES (SHEEP). //P024
 41
 F-NT2RP2001394//POLYHOMEOTIC-PROXIMAL CHROMATIN PR
 OTEIN. //0.024:39:53//DROSOPHILA MELANOGASTER (FRUI
 T FLY). //P39769
 F-NT2RP2001397//G2/MITOTIC-SPECIFIC CYCLIN B2. //1.
 4e-46:125:78//MESOCRICETUS AURATUS (GOLDEN HAMSTE
 R). //P37883
 F-NT2RP2001420//SALIVARY PROLINE-RICH PROTEIN PREC
 URSOR (CLONES CP3, CP4AND CP5) [CONTAINS: BASIC PE
 PTIDE IB-6; PEPTIDE P-H]. //0.00018:113:38//HOMO SA
 PIENS (HUMAN). //P04280
 F-NT2RP2001423//HYPOTHETICAL 9.4 KD PROTEIN IN GP3
 1-CD INTERGENIC REGION(ORF A). //0.90:23:43//BACTER
 IOPHAGE T4. //P17307
 F-NT2RP2001427//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/2.2e-11:38:68//HOMO SAPIENS (HUMAN). //P3918
 8
 F-NT2RP2001436//DYNEIN LIGHT INTERMEDIATE CHAIN 2,
 CYTOSOLIC (LIC53/55) (LIC-2). //0.25:124:28//RATTUS
 NORVEGICUS (RAT). //Q62698
 F-NT2RP2001440//14-3-3 PROTEIN GAMMA (PROTEIN KINA
 SE C INHIBITOR PROTEIN-1) (KCIP-1). //4.8e-62:145:9
 0//RATTUS NORVEGICUS (RAT). //P35214
 F-NT2RP2001445
 F-NT2RP2001449//CLEAVAGE AND POLYADENYLATION SPECI
 FICITY FACTOR, 100 KDSUBUNIT (CPSF 100 KD SUBUNI
 T). //9.5e-118:226:95//BOS TAURUS (BOVINE). //Q10568 50 8
- F-NT2RP2001450
 F-NT2RP2001467//SHORT NEUROTOXIN 1 (TOXIN V-II-1).
 //1.0:25:40//BUNGARUSFASCIATUS (BANDED KRAIT). //P1
 0808
 F-NT2RP2001506
 F-NT2RP2001511//HYPOTHETICAL 115.4 KD PROTEIN ZK75
 7.3 IN CHROMOSOME III. //0.49:124:29//CAENORHABDITI
 S ELEGANS. //P34681
 F-NT2RP2001520//VITAMIN D-DEPENDENT CALCIUM-BINDIN
 10 G PROTEIN, INTESTINAL(CABP) (CALBINDIN D9K). //0.03
 5:71:33//HOMO SAPIENS (HUMAN). //P29377
 F-NT2RP2001526
 F-NT2RP2001536//METALLOTHIONEIN-I (MT-1). //1.0:19:
 42//COLUMBA LIVIA (DOMESTIC PIGEON). //P15786
 F-NT2RP2001560//CUTICLE COLLAGEN 12 PRECURSOR. //0.
 0018:144:35//CAENORHABDITIS ELEGANS. //P20630
 F-NT2RP2001569//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!/2.7e-31:102:67//HOMO SAPIENS (HUMAN). //P39
 194
 F-NT2RP2001576//SMP3 PROTEIN. //0.00016:75:36//SACC
 HAROMYCES CEREVISIAE (BAKER'S YEAST). //Q04174
 F-NT2RP2001581//TRANSMEMBRANE PROTEIN SEX PRECURSO
 R. //0.040:46:36//HOMOSAPIENS (HUMAN). //P51805
 F-NT2RP2001597//PROBABLE E4 PROTEIN. //0.00042:113:
 34//HUMAN PAPILOMAVIRUS TYPE 5. //P06924
 F-NT2RP2001601
 F-NT2RP2001613//HOMEBOX PROTEIN SAX-1 (CHOX-3) (F
 RAGMENT). //0.14:59:32//GALLUS GALLUS (CHICKEN). //P
 19601
 30 F-NT2RP2001628//A-AGGLUTININ ATTACHMENT SUBUNIT PR
 ECURSOR. //0.056:140:33//SACCHAROMYCES CEREVISIAE
 (BAKER'S YEAST). //P32323
 F-NT2RP2001634//ALPHA-CATENIN. //7.1e-12:152:35//DR
 OSOPHILA MELANOGASTER(FRUIT FLY). //P35220
 F-NT2RP2001660//HYPOTHETICAL 80.4 KD PROTEIN IN SM
 C3-MRPL8 INTERGENIC REGION. //0.43:119:26//SACCHARO
 MYCES CEREVISIAE (BAKER'S YEAST). //P40358
 F-NT2RP2001663//ALPHA ENOLASE (EC 4.2.1.11) (2-PHO
 SPHO-D-GLYCERATE HYDRO-LYASE) (NON- NEURAL ENOLAS
 40 E) (NNE) (PHOSPHOPYRUVATE HYDRATASE). //1.2e-26:12
 6:56//HOMO SAPIENS (HUMAN). //P06733
 F-NT2RP2001675//HYPOTHETICAL 107.7 KD PROTEIN IN R
 PSO 5' REGION (ORF1). //0.25:148:25//CAMPYLOBACTER J
 EJUNI. //Q46089
 F-NT2RP2001677//ACIDIC PROLINE-RICH PROTEIN PRECUR
 SOR (CLONE PRP25) (FRAGMENT). //0.010:101:31//RATTU
 S NORVEGICUS (RAT). //P10164
 F-NT2RP2001678//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/2.6e-18:83:61//HOMO SAPIENS (HUMAN). //P3918

3003

F-NT2RP2001699//PROTEIN C14. //0.98:51:31//VACCINIA VIRUS (STRAIN COPENHAGEN). //P21045
 F-NT2RP2001720//MEROZOITE SURFACE ANTIGEN 2 PRECURSOR (MSA-2) (ALLELIC FORM 1). //0.16:145:30//PLASMODIUM FALCIPARUM (ISOLATE CAMP / MALAYSIA). //Q99317
 F-NT2RP2001721//MALE-SPECIFIC LETHAL-2 PROTEIN. //0.00090:48:39//DROSOPHILA MELANOGASTER (FRUIT FL Y). //P50534
 F-NT2RP2001740//ANNEXIN VII (SYNEXIN) (FRAGMENT). //0.50:43:25//BOS TAURUS (BOVINE). //P20072
 F-NT2RP2001748//SALIVARY PROLINE-RICH PROTEIN PO (ALLELE K) [CONTAINS: PEPTIDE P-D] (FRAGMENT). //0.77:111:28//HOMO SAPIENS (HUMAN). //P10162
 F-NT2RP2001762
 F-NT2RP2001813//PHOTOSYSTEM I REACTION CENTRE SUBUNIT VIII (PSI-I). //1.0:22:40//PICEA ABIES (NORWAY SPRUCE) (PICEA EXCELSA). //047040
 F-NT2RP2001839//SCY1 PROTEIN. //6.8e-17:204:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53009
 F-NT2RP2001861//D15KZ1 PROTEIN (FRAGMENT). //0.31:56:39//MUS MUSCULUS (MOUSE). //Q61466
 F-NT2RP2001869//CORNEODESMOSIN (S PROTEIN) (FRAGMENT). //0.97:78:30//SUSSCROFA (PIG). //019084
 F-NT2RP2001876//ALLOGRAFT INFLAMMATORY FACTOR-1 (AIF-1) (IONIZED CALCIUM-BINDING ADAPTER MOLECULE 1). //3.5e-36:106:66//HOMO SAPIENS (HUMAN). //P55008
 F-NT2RP2001883//CATHEPSIN L (EC 3.4.22.15). //0.95:29:41//OVIS ARIES (SHEEP). //Q10991
 F-NT2RP2001898//TYPE II INOSITOL-1,4,5-TRISPHOSPHATE 5-PHOSPHATASE PRECURSOR (EC 3.1.3.56) (5PTASE) (FRAGMENT). //1.6e-84:185:88//HOMO SAPIENS (HUMAN). //P32019
 F-NT2RP2001900//ACTIN-LIKE PROTEIN ARP5. //1.1e-17:180:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53946
 F-NT2RP2001907//HYPHAL WALL PROTEIN 1 (CELL ELONGATION PROTEIN 2). //0.13:108:27//CANDIDA ALBICANS (YEAST). //P46593
 F-NT2RP2001926//HYPOTHETICAL 7.6 KD PROTEIN YCF33. //0.55:57:26//CYANOPHORA PARADOXA. //P48273
 F-NT2RP2001936
 F-NT2RP2001943//HYPOTHETICAL 57.7 KD PROTEIN IN AIP1-CTF13 INTERGENIC REGION. //1.8e-13:208:22//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q04305
 F-NT2RP2001946//HYPOTHETICAL 13.0 KD PROTEIN IN ALGR3 3' REGION. //0.59:76:28//PSEUDOMONAS AERUGINOSA. //P21485
 F-NT2RP2001947//ZINC FINGER PROTEIN DAN (NO3). //0.53:68:29//RATTUS NORVEGICUS (RAT). //Q06880
 F-NT2RP2001969//CHLOROPLAST 30S RIBOSOMAL PROTEIN

3004

S18. //0.0015:52:34//CHLORELLA VULGARIS. //P56353
 F-NT2RP2001976//DILUTE MYOSIN HEAVY CHAIN, NON-MUSCLE (MYOSIN 5A). //9.5e-07:201:22//MUS MUSCULUS (MOUSE). //Q99104
 F-NT2RP2001985//PROLINE-RICH PROTEIN MP-2 PRECURSOR. //0.016:90:32//MUS MUSCULUS (MOUSE). //P05142
 F-NT2RP2001991//SODIUM- AND CHLORIDE-DEPENDENT TRANSPORTER NTT73. //8.0e-14:47:76//RATTUS NORVEGICUS (RAT). //Q08469
 10 F-NT2RP2002025//NG-CAM RELATED CELL ADHESION MOLECULE PRECURSOR (NR-CAM) (BRAVO). //2.9e-30:211:42//GALLUS GALLUS (CHICKEN). //P35331
 F-NT2RP2002032//FLOCCULANT-ACTIVE PROTEINS M02.1 AND M02.2. //0.23:20:40//MORINGA OLEIFERA (HORSE RADISH TREE) (MORINGA PTERYGOSPERMA). //P24303
 F-NT2RP2002033//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!! //0.88:27:62//HOMO SAPIENS (HUMAN). //P39193
 F-NT2RP2002041
 F-NT2RP2002046//MATING PROCESS PROTEIN MID2 (SERINE-RICH PROTEIN SMS1) (PROTEIN KINASE A INTERFERENCE PROTEIN). //1.0:85:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P36027
 F-NT2RP2002047
 F-NT2RP2002058//DOM34 INTERACTING PROTEIN 2. //9.4e-25:165:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //Q12220
 F-NT2RP2002066//TIGHT JUNCTION PROTEIN ZO-1 (TIGHT JUNCTION PROTEIN 1). //5.7e-12:108:41//HOMO SAPIENS (HUMAN). //Q07157
 30 F-NT2RP2002070//CYTOCHROME C OXIDASE POLYPEPTIDE I (EC 1.9.3.1) (FRAGMENT). //0.88:28:50//ASTERINA PECTINIFERA (STARFISH). //P11958
 F-NT2RP2002076//TRP-ASP REPEATS CONTAINING PROTEIN RBA-2. //0.0031:124:27//CAENORHABDITIS ELEGANS. //P90916
 F-NT2RP2002078//KERATIN, GLYCINE/TYROSINE-RICH OF HAIR. //0.82:30:40//OVIS ARIES (SHEEP). //Q02958
 F-NT2RP2002079//OUTER DENSE FIBER PROTEIN. //0.34:41:39//HOMO SAPIENS (HUMAN). //Q14990
 40 F-NT2RP2002099//HETEROGENOUS NUCLEAR RIBONUCLEOPROTEIN U (HNRNP U). //5.2e-08:81:48//HOMO SAPIENS (HUMAN). //Q00839
 F-NT2RP2002105//COLLAGEN 1(X) CHAIN PRECURSOR. //0.0012:100:34//BOS TAURUS (BOVINE). //P23206
 F-NT2RP2002124//EARLY GROWTH RESPONSE PROTEIN 1 (EGR-1) (KROX24) (TRANSCRIPTION FACTOR ETR103) (ZINC FINGER PROTEIN 225) (AT225). //0.74:72:31//HOMO SAPIENS (HUMAN). //P18146
 F-NT2RP2002137//NEUROTOXIN B-II. //1.0:27:44//CEREBRATULUS LACTEUS (MILKYRIBBON WORM). //P01526

3005

F-NT2RP2002154//GALECTIN-3 (GALACTOSE-SPECIFIC LECTIN 3) (MAC-2 ANTIGEN) (IGE-BINDING PROTEIN) (35 KD LECTIN) (CARBOHYDRATE BINDING PROTEIN 35) (CBP 35) (LAMININ-BINDING PROTEIN) (LECTIN L-29) (L-34 GALACTOSIDE-BINDING LECTIN). //0.0029:112:34//MUS MUSCULUS (MOUSE). //P16110
 F-NT2RP2002172
 F-NT2RP2002185//UBIQUITIN-LIKE PROTEIN DSK2. //1.8e-07:87:40//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P48510
 F-NT2RP2002192
 F-NT2RP2002193//CUTICLE COLLAGEN 40. //0.0062:70:37//CAENORHABDITIS ELEGANS. //P34804
 F-NT2RP2002208//PEROXISOME ASSEMBLY PROTEIN PEX10 (PEROXIN-10). //0.00011:45:40//HOMO SAPIENS (HUMAN). //060683
 F-NT2RP2002219
 F-NT2RP2002231//V-TYPE SODIUM ATP SYNTHASE SUBUNIT E (EC 3.6.1.34) (NA(+)- TRANSLOCATING ATPASE SUBUNIT E). //1.0:68:32//ENTEROCOCCUS HIRAE. //P43436
 F-NT2RP2002235//INFECTED CELL PROTEIN ICP34.5 (NEUROVIRULENCE FACTOR ICP34.5). //0.0022:66:45//HERPES SIMPLEX VIRUS (TYPE 1 / STRAIN CVG-2). //P37318
 F-NT2RP2002252//DNA-DIRECTED RNA POLYMERASE II LARGE SUBUNIT (EC 2.7.7.6) (RPB1) (FRAGMENT). //0.071:110:31//CRICETULUS GRISEUS (CHINESE HAMSTER). //P11414
 F-NT2RP2002256//CYTOCHROME P450 26 (EC 1.14.-.-) (RETINOIC ACID-METABOLIZING CYTOCHROME) (P450RA1) (RETINOIC ACID 4-HYDROXYLASE). //3.1e-31:75:84//MUS MUSCULUS (MOUSE). //055127
 F-NT2RP2002259//L-MYC-1 PROTO-ONCOGENE PROTEIN. //1.9e-17:41:90//HOMO SAPIENS (HUMAN). //P12524
 F-NT2RP2002270//HYPOTHETICAL 26.0 KD PROTEIN IN CYB5-LEU4 INTERGENIC REGION. //2.1e-27:164:36//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53930
 F-NT2RP2002292//IMMEDIATE-EARLY PROTEIN RSP40. //0.018:107:23//PSEUDORABIES VIRUS (STRAIN KAPLAN) (PRV). //P24827
 F-NT2RP2002312//PHOSPHATIDATE CYTIDYLTRANSFERASE (EC 2.7.7.41) (CDP-DIGLYCERIDE SYNTHETASE) (CDP-DIGLYCERIDE PYROPHOSPHORYLASE) (CDP-DIACYLGLYCEROL SYNTHASE) (CDS) (CTP:PHOSPHATIDATE CYTIDYLTRANSFERASE) (CDP-DAGSYNTHASE). //1.4e-52:174:55//HOMO SAPIENS (HUMAN). //Q92903
 F-NT2RP2002316//HISTONE H1.C6/H1.C9. //1.0:40:40//TRYPANOSOMA CRUZI. //P40269
 F-NT2RP2002325//PEROXISOMAL MEMBRANE PROTEIN PMP30A (PMP31) (PEROXIN-11A). //2.2e-06:145:26//CANDIDA BOIDINII (YEAST). //Q00316

3006

F-NT2RP2002333//HYPOTHETICAL 39.1 KD PROTEIN IN RNPB-SOHA INTERGENIC REGION (ORF 3). //0.30:86:32//ESCHERICHIA COLI. //P23524
 F-NT2RP2002373//SYNAPSINS IA AND IB. //0.080:145:31//BOS TAURUS (BOVINE). //P17599
 F-NT2RP2002385//ENV POLYPROTEIN PRECURSOR (COAT POLYPROTEIN) [CONTAINS: KNOB PROTEIN GP70; SPIKE PROTEIN P15E; R PROTEIN]. //0.021:66:28//MINK CELL FOCUS-FORMING MURINE LEUKEMIA VIRUS (ISOLATE CI-3). //P03388
 F-NT2RP2002394
 F-NT2RP2002408//COLLAGEN ALPHA 1(I) CHAIN (FRAGMENTS). //0.00030:107:37//BOS TAURUS (BOVINE). //P02453
 F-NT2RP2002426
 F-NT2RP2002439//CIRCUMSPOROZOITE PROTEIN PRECURSOR (CS). //0.00032:79:32//PLASMODIUM BERGHEI (STRAIN ANKA). //P23093
 F-NT2RP2002442//HESA PROTEIN. //6.0e-16:163:30//PLECTONEMA BORYANUM. //P46037
 F-NT2RP2002457
 F-NT2RP2002464//HYPOTHETICAL 60.7 KD PROTEIN C56F8.17C IN CHROMOSOME I. //9.3e-18:165:32//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10264
 F-NT2RP2002475//CYSTEINE-RICH HEART PROTEIN (HCRHP). //0.91:45:35//HOMO SAPIENS (HUMAN). //P50238
 F-NT2RP2002479//ATP-BINDING CASSETTE TRANSPORTER 7 PRECURSOR (ABC TRANSPORTER 7 PROTEIN). //6.8e-96:186:94//HOMO SAPIENS (HUMAN). //075027
 F-NT2RP2002498//HYPOTHETICAL MERCURIC RESISTANCE PROTEIN MERC. //0.65:37:45//PSEUDOMONAS AERUGINOSA. //P04139
 F-NT2RP2002503//ZINC FINGER PROTEIN 45 (BRC1744). //1.3e-31:124:59//HOMOSAPIENS (HUMAN). //Q02386
 F-NT2RP2002504//NUCLEAR PORE COMPLEX PROTEIN NUP155 (NUCLEOPORIN NUP155) (155 KD NUCLEOPORIN) (P140). //1.2e-123:240:92//RATTUS NORVEGICUS (RAT). //P37199
 F-NT2RP2002520//ACIDIC PROLINE-RICH PROTEIN HP43A PRECURSOR. //0.94:83:28//MESOCRICETUS AURATUS (GOLDEN HAMSTER). //P06680
 F-NT2RP2002537//HYPOTHETICAL 55.1 KD PROTEIN B0416.5 IN CHROMOSOME X. //4.0e-10:194:23//CAENORHABDITIS ELEGANS. //Q11073
 F-NT2RP2002546
 F-NT2RP2002549//G2/MITOTIC-SPECIFIC CYCLIN C13-1 (A-LIKE CYCLIN) (FRAGMENT). //0.98:65:30//DAUCUS CAROTA (CARROT). //P25010
 F-NT2RP2002591//ZINC FINGER PROTEIN 84 (ZINC FINGER PROTEIN HPF2). //2.6e-19:60:61//HOMO SAPIENS (HUMAN). //P51523

3007

F-NT2RP2002595//ANNEXIN VII (SYNEXIN). //1.2e-15:12
 1:49//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //Q9212
 5
 F-NT2RP2002606//PROTEIN TRANSPORT PROTEIN SEC2. //
 0.00034:98:31//SACCHAROMYCES CEREVISIAE (BAKER'S Y
 EAST). //P17065
 F-NT2RP2002609//HYPOTHETICAL 52.0 KD PROTEIN IN CL
 B6-SPT6 INTERGENIC REGION. //0.00022:79:39//SACCHAR
 OMYCES CEREVISIAE (BAKER'S YEAST). //P53264
 F-NT2RP2002618//PROTEIN ARGININE N-METHYLTRANSFERA 10
 SE 1 (EC 2.1.1.-). //6.2e-37:180:44//RATTUS NORVEGI
 CUS (RAT). //Q63009
 F-NT2RP2002621//NADH-UBIQUINONE OXIDOREDUCTASE CHA
 IN 5 (EC 1.6.5.3) (FRAGMENT). //0.98:37:35//LEMUR C
 ATTA (RING-TAILED LEMUR). //Q34879
 F-NT2RP2002643//INFECTED CELL PROTEIN ICP34.5 (NEU
 ROVIRULENCE FACTOR ICP34.5). //0.042:77:32//HERPES
 SIMPLEX VIRUS (TYPE 1 / STRAIN MGH-10). //P37319
 F-NT2RP2002672//PROTEIN Q300. //0.0018:41:43//MUS M
 USCULUS (MOUSE). //Q02722
 F-NT2RP2002701//HYPOTHETICAL 72.5 KD PROTEIN C2F7.
 10 IN CHROMOSOME I. //3.6e-17:100:42//SCHIZOSACCHAR
 OMYCES POMBE (FISSION YEAST). //Q09701
 F-NT2RP2002706//IMMEDIATE-EARLY PROTEIN IE180. //0.
 00027:139:33//PSEUDORABIES VIRUS (STRAIN KAPLAN)
 (PRV). //P33479
 F-NT2RP2002710//SH3-BINDING PROTEIN 3BP-1. //6.9e-0
 9:96:40//MUS MUSCULUS(MOUSE). //P55194
 F-NT2RP2002727//TUBERIN (TUBEROUS SCLEROSIS 2 HOMO
 LOG PROTEIN). //3.6e-20:160:36//RATTUS NORVEGICUS 30
 (RAT). //P49816
 F-NT2RP2002736
 F-NT2RP2002740
 F-NT2RP2002741//RH01 GDP-GTP EXCHANGE PROTEIN 2. //
 2.0e-07:178:28//SACCHAROMYCES CEREVISIAE (BAKER'S
 YEAST). //P51862
 F-NT2RP2002750//!!!! ALU SUBFAMILY SB2 WARNING ENT
 RY !!!!!//1.6e-09:43:72//HOMO SAPIENS (HUMAN). //P39
 191
 F-NT2RP2002752//LOW CALCIUM RESPONSE LOCUS PROTEIN 40
 T. //0.95:33:39//YERSINIA PSEUDOTUBERCULOSIS. //Q00
 932
 F-NT2RP2002753//ENDOGLUCANASE EG-1 PRECURSOR (EC
 3.2.1.4) (ENDO-1,4-BETA-GLUCANASE) (CELLULASE). //
 0.71:78:33//TRICHERDIA LONGIBRACHIATUM. //Q12714
 F-NT2RP2002769//50 KD SPICULE MATRIX PROTEIN PRECU
 RSOR. //0.44:76:32//STRONGYLOCENTROTUS PURPURATUS
 (PURPLE SEA URCHIN). //P11994
 F-NT2RP2002778
 F-NT2RP2002800//CRAMBIN. //0.99:20:50//CRAMBE ABYSS 50

3008

INICA (ABYSSINIAN CRAMBE). //P01542
 F-NT2RP2002839//SALIVARY PROLINE-RICH PROTEIN PREC
 URSOR (CLONE CP7) [CONTAINS: BASIC PEPTIDE P-F] (F
 RAGMENT). //0.010:87:31//HOMO SAPIENS (HUMAN). //P02
 812
 F-NT2RP2002857//ACIDIC PROLINE-RICH PROTEIN PRECUR
 SOR (CLONE PRP33). //0.00018:57:45//RATTUS NORVEGIC
 US (RAT). //P04474
 F-NT2RP2002862//HYPOTHETICAL 27.1 KD PROTEIN UFD4-
 CAP1 INTERGENIC REGION. //7.2e-27:140:40//SACCHAROM
 YCES CEREVISIAE (BAKER'S YEAST). //P33201
 F-NT2RP2002880//DNA REPAIR PROTEIN RAD32. //0.83:6
 7:28//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //
 Q09683
 F-NT2RP2002891//HOMEBOX PROTEIN DLX-2 (DLX-5) (FR
 AGMENT). //0.99:70:24//RATTUS NORVEGICUS (RAT). //Q6
 4204
 F-NT2RP2002925//ALPHA-1D ADRENERGIC RECEPTOR (ALPH
 A 1D-ADRENOCEPTOR) (ALPHA-1A ADRENERGIC RECEPTOR).
 20 //0.31:48:43//HOMO SAPIENS (HUMAN). //P25100
 F-NT2RP2002928//CELL DIVISION CONTROL PROTEIN 40. /
 /2.8e-26:142:42//SACCHAROMYCES CEREVISIAE (BAKER'S
 YEAST). //P40968
 F-NT2RP2002929//HYPOTHETICAL 46.2 KD TRP-ASP REPEA
 TS CONTAINING PROTEIN D2013.2 IN CHROMOSOME II. //2.
 0e-31:186:35//CAENORHABDITIS ELEGANS. //Q18964
 F-NT2RP2002939//ADENYLATE CYCLASE, TYPE V (EC 4.6.
 1.1) (ATP PYROPHOSPHATE-LYASE) (CA(2+)-INHIBITABLE
 ADENYLYL CYCLASE). //0.0022:98:39//CANIS FAMILIARI
 S (DOG). //P30803
 F-NT2RP2002954//U2 SMALL NUCLEAR RIBONUCLEOPROTEIN
 A' (U2 SNRNP-A'). //0.0019:107:30//ARABIDOPSIS THA
 LIANA (MOUSE-EAR CRESS). //P43333
 F-NT2RP2002959//UBIQUITIN-CONJUGATING ENZYME E2-17
 KD 2 (EC 6.3.2.19) (UBIQUITIN- PROTEIN LIGASE) (U
 BIQUITIN CARRIER PROTEIN) (E2(17)KB 2). //2.8e-11:3
 3:81//HOMO SAPIENS (HUMAN), MUS MUSCULUS (MOUSE),
 RATTUS NORVEGICUS (RAT), AND XENOPUS LAEVIS (AFRIC
 AN CLAWED FROG). //P51669
 F-NT2RP2002979
 F-NT2RP2002980//30S RIBOSOMAL PROTEIN S10. //1.1e-0
 9:98:36//MYCOPLASMA CAPRICOLUM. //P10129
 F-NT2RP2002986//RING CANAL PROTEIN (KELCH PROTEI
 N). //1.1e-19:141:39//DROSOPHILA MELANOGASTER (FRUI
 T FLY). //Q04652
 F-NT2RP2002987//!!!! ALU SUBFAMILY SC WARNING ENTR
 Y !!!!!//1.3e-07:78:47//HOMO SAPIENS (HUMAN). //P391
 92
 F-NT2RP2002993//DNA-DIRECTED RNA POLYMERASE I 135
 KD POLYPEPTIDE (EC 2.7.7.6) (RNA POLYMERASE I SUBU

3009

NIT 2) (RPA135) (RNA POLYMERASE I 127 KD SUBUNIT).
 //8.0e-77:165:85//RATTUS NORVEGICUS (RAT).//054888
 F-NT2RP2003000//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//2.8e-19:62:64//HOMO SAPIENS (HUMAN).//P391
 94
 F-NT2RP2003034//HYPOTHETICAL PROTEIN HI1458.//1.0:
 42:35//HAEMOPHILUS INFLUENZAE.//P44204
 F-NT2RP2003073//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!//0.0051:16:87//HOMO SAPIENS (HUMAN).//P3918
 9
 F-NT2RP2003099
 F-NT2RP2003108//BASIC PROLINE-RICH PEPTIDE IB-1.//
 0.84:47:34//HOMO SAPIENS (HUMAN).//P04281
 F-NT2RP2003117
 F-NT2RP2003121//HYPOTHETICAL 96.7 KD PROTEIN IN ST
 E2-FRS2 INTERGENIC REGION.//9.0e-08:99:29//SACCHAR
 OMYCES CEREVISIAE (BAKER'S YEAST).//P43572
 F-NT2RP2003125//TRANSCRIPTION REGULATOR PROTEIN BA
 CH2 (BTB AND CNC HOMOLOG 2).//9.2e-08:134:28//MUS
 MUSCULUS (MOUSE).//P97303
 F-NT2RP2003129
 F-NT2RP2003137//UBIQUITIN.//3.4e-06:70:30//NEUROSP
 ORA CRASSA.//P13117
 F-NT2RP2003157//HYPOTHETICAL 37.0 KD PROTEIN B049
 5.8 IN CHROMOSOME II.//7.8e-13:84:40//CAENORHABDI
 TIS ELEGANS.//Q09217
 F-NT2RP2003158//26S PROTEASOME REGULATORY SUBUNIT
 S3 (PROTEASOME SUBUNIT P58).//3.1e-65:155:84//HOMO
 SAPIENS (HUMAN).//043242
 F-NT2RP2003161//PROLINE-RICH PROTEIN MP-2 PRECURSO
 R.//0.0011:59:42//MUS MUSCULUS (MOUSE).//P05142
 F-NT2RP2003164//ZYXIN.//0.0037:85:36//MUS MUSCULUS
 (MOUSE).//Q62523
 F-NT2RP2003165//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//1.2e-24:77:64//HOMO SAPIENS (HUMAN).//P391
 94
 F-NT2RP2003177//MALE SPECIFIC SPERM PROTEIN MST84D
 B.//0.55:38:39//DROSOPHILA MELANOGASTER (FRUIT FL
 Y).//Q01643
 F-NT2RP2003194//HYPOTHETICAL 12.5 KD PROTEIN ZK63
 7.2 IN CHROMOSOME III.//2.3e-14:87:37//CAENORHABDI
 TIS ELEGANS.//P30629
 F-NT2RP2003206//NADH-UBIQUINONE OXIDOREDUCTASE CHA
 IN 3 (EC 1.6.5.3).//1.0:100:28//DIDELPHIS MARSUPIA
 LIS VIRGINIANA (NORTH AMERICAN OPOSSUM).//P41306
 F-NT2RP2003228//DNA REPLICATION LICENSING FACTOR M
 CM4 (CDC21 HOMOLOG) (P1-CDC21).//9.3e-82:211:81//H
 OMO SAPIENS (HUMAN).//P33991
 F-NT2RP2003230//SEC14 CYTOSOLIC FACTOR (PHOSPHATID
 YLINOSITOL/PHOSPHATIDYLCHOLINE TRANSFER PROTEIN)

3010

(PI/PC TP).//1.0:51:31//CANDIDA GLABRATA (YEAST)
 (TORULOPSIS GLABRATA).//P53989
 F-NT2RP2003237//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!//5.1e-44:66:84//HOMO SAPIENS (HUMAN).//P391
 94
 F-NT2RP2003243//M PROTEIN, SEROTYPE 5 PRECURSOR.//
 0.027:204:23//STREPTOCOCCUS PYOGENES.//P02977
 F-NT2RP2003265//BP4A PROTEIN.//0.95:35:34//BRASSIC
 A NAPUS (RAPE).//P41505
 10 F-NT2RP2003272//ANTER-SPECIFIC PROLINE-RICH PROTEI
 N APG (PROTEIN CEX) (FRAGMENT).//5.5e-06:78:35//BR
 ASSICA NAPUS (RAPE).//P40603
 F-NT2RP2003277//NAM7 PROTEIN (NONSENSE-MEDIATED MR
 NA DECAY PROTEIN 1) (UP-FRAMESHIFT SUPPRESSOR 1).//
 1.9e-19:145:43//SACCHAROMYCES CEREVISIAE (BAKER'S
 YEAST).//P30771
 F-NT2RP2003280
 F-NT2RP2003286//RNA 3'-TERMINAL PHOSPHATE CYCLASE
 (EC 6.5.1.4) (RNA-3'-PHOSPHATE CYCLASE) (RNA CYCLA
 SE).//2.1e-32:137:42//SACCHAROMYCES CEREVISIAE (BA
 KER'S YEAST).//Q08096
 20 F-NT2RP2003293//ZINC FINGER PROTEIN 83 (ZINC FINGE
 R PROTEIN HPF1).//7.7e-12:175:33//HOMO SAPIENS (HU
 MAN).//P51522
 F-NT2RP2003295//PTB-ASSOCIATED SPLICING FACTOR (PS
 F).//0.055:44:45//HOMOSAPIENS (HUMAN).//P23246
 F-NT2RP2003297
 F-NT2RP2003307//KINESIN LIGHT CHAIN (KLC).//2.0e-1
 8:87:49//RATTUS NORVEGICUS (RAT).//P37285
 30 F-NT2RP2003308//CROOKED NECK PROTEIN.//2.1e-91:24
 4:67//DROSOPHILA MELANOGASTER (FRUIT FLY).//P17886
 F-NT2RP2003329//HYPOTHETICAL 54.9 KD PROTEIN C02F
 5.7 IN CHROMOSOME III.//5.8e-57:186:55//CAENORHABD
 ITIS ELEGANS.//P34284
 F-NT2RP2003339//SHORT NEUROTOXIN 1 (NEUROTOXIN ALP
 HA).//0.98:11:72//DENDROASPIS POLYLEPIS POLYLEPIS
 (BLACK MAMBA).//P01416
 F-NT2RP2003347//60S RIBOSOMAL PROTEIN L38.//0.83:4
 2:33//OSTERTAGIA OSTERTAGI.//061570
 40 F-NT2RP2003367//SYNERGISTIC-TYPE VENOM PROTEIN C9S
 3, CHAIN 1.//1.0:37:35//DENDROASPIS ANGUSTICEPS (E
 ASTERN GREEN MAMBA).//P01408
 F-NT2RP2003391//MRNA TRANSPORT REGULATOR MTR10.//
 3.3e-11:229:24//SACCHAROMYCES CEREVISIAE (BAKER'S
 YEAST).//Q99189
 F-NT2RP2003393//PROTOCHLOROPHYLLIDE REDUCTASE CHLB
 SUBUNIT (EC 1.3.1.33) (NADPH- PROTOCHLOROPHYLLIDE
 OXIDOREDUCTASE CHLB SUBUNIT) (FRAGMENT).//0.94:29:
 34//ARAUCARIA HETEROPHYLLA.//P37843
 50 F-NT2RP2003394

3011

F-NT2RP2003401//60 KD CHAPERONIN (PROTEIN CPN60) (GROEL PROTEIN).//0.95:125:28//THERMUS AQUATICUS (SUBSP. THERMOPHILUS).//P45746
 F-NT2RP2003433//PROTEIN TRANSPORT PROTEIN SEC61 ALPHA SUBUNIT.//9.8e-78:178:84//RATTUS NORVEGICUS (RAT).//P38378
 F-NT2RP2003445
 F-NT2RP2003446//HYPOTHETICAL PROTEIN E-115.//0.00030:106:33//HUMAN ADENOVIRUS TYPE 2.//P03290
 F-NT2RP2003456//PHOTOSYSTEM II REACTION CENTRE M P ROEIN.//1.0:27:51//MARCHANTIA POLYMORPHA (LIVERWORT).//P12168
 F-NT2RP2003466//LINOLEOYL-COA DESATURASE (EC 1.14.99.25) (DELTA(6)-DESATURASE).//6.7e-06:108:32//SYNCHOCYSTIS SP. (STRAIN PCC 6803).//Q08871
 F-NT2RP2003480//TRANSCRIPTION FACTOR BF-2 (BRAIN FACTOR 2) (BF2) (CBF-2) (T-14-6).//7.2e-15:38:50//GALLUS GALLUS (CHICKEN).//Q98937
 F-NT2RP2003499//SE5 ANTIGEN.//0.090:114:32//RATTUS NORVEGICUS (RAT).//Q63003
 F-NT2RP2003506//NADPH-CYTOCHROME P450 REDUCTASE (EC 1.6.2.4) (CPR).//2.0e-11:91:43//SUS SCROFA (PIG).//P04175
 F-NT2RP2003511//PARMYOSIN, SHORT FORM (MIMIPARMYOSIN).//0.0020:108:25//DROSOPHILA MELANOGASTER (FRUIT FLY).//P35416
 F-NT2RP2003513//PTB-ASSOCIATED SPLICING FACTOR (PSF).//1.2e-05:96:36//HOMO SAPIENS (HUMAN).//P23246
 F-NT2RP2003517//HYPOTHETICAL 12.9 KD PROTEIN CY49.27.//0.0059:22:31//MYCOBACTERIUM TUBERCULOSIS.//Q10696
 F-NT2RP2003522//HYPOTHETICAL 10.0 KD PROTEIN.//1.0:65:30//THERMOPROTEUSTENAX VIRUS 1 (STRAIN KRA1) (TTV1).//P19283
 F-NT2RP2003533//LINE-1 REVERSE TRANSCRIPTASE HOMOLOG.//8.7e-18:94:54//HOMO SAPIENS (HUMAN).//P08547
 F-NT2RP2003543//SYNAPSINS IA AND IB.//0.045:101:35//RATTUS NORVEGICUS (RAT).//P09951
 F-NT2RP2003559//ITBA2 PROTEIN (DXS9879E).//0.98:37:37//HOMO SAPIENS (HUMAN).//Q14657
 F-NT2RP2003564//52 KD RO PROTEIN (SJOEGREN SYNDROME TYPE A ANTIGEN (SS-A)) (RO(SS-A)).//6.4e-35:175:44//HOMO SAPIENS (HUMAN).//P19474
 F-NT2RP2003567//HYPOTHETICAL 11.2 KD PROTEIN T18D 3.7 IN CHROMOSOME X.//0.72:82:34//CAENORHABDITIS ELEGANS.//Q22544
 F-NT2RP2003581//HOMEBOX PROTEIN OTX1.//0.90:61:37//MUS MUSCULUS (MOUSE).//P80205
 F-NT2RP2003596//ELONGATION FACTOR P (EF-P).//0.83:61:32//MYCOPLASMA GENITALIUM.//P47272

3012

F-NT2RP2003604//ALPHA-CATENIN.//1.5e-11:152:33//DROSOPHILA MELANOGASTER (FRUIT FLY).//P35220
 F-NT2RP2003629//PHOSPHOLIPASE A2 ALPHA (EC 3.1.1.4) (PHOSPHATIDYLCHOLINE2-ACYLHYDROLASE).//0.97:85:27//CROTALUS ADAMANTEUS (EASTERN DIAMONDBACKRATTLE SNAKE).//P00623
 F-NT2RP2003643//ACYLNEURAMINATE CYTIDYLYLTRANSFERASE (EC 2.7.7.43) (CMP-N-ACETYLNEURAMINIC ACID SYNTHETASE) (CMP-NEUNAC SYNTHETASE) (CMP-SIALICACID SYNTHETASE).//3.9e-12:84:40//NEISSERIA MENINGITIDIS.//Q57385
 F-NT2RP2003668//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//5.0e-33:74:81//HOMO SAPIENS (HUMAN).//P39195
 F-NT2RP2003687//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//5.7e-05:40:67//HOMO SAPIENS (HUMAN).//P39188
 F-NT2RP2003691//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!!//2.5e-37:56:67//HOMO SAPIENS (HUMAN).//P39194
 F-NT2RP2003702//HYPOTHETICAL OXIDOREDUCTASE IN INLA 5' REGION (EC 1.-.-) (ORFA).//1.3e-07:98:37//LISTERIA MONOCYTOGENES.//P25145
 F-NT2RP2003704//GAMMA-GLUTAMYLTRANSPEPTIDASE 5 PRECURSOR (EC 2.3.2.2) (GAMMA-GLUTAMYLTRANSFERASE 5) (GGT-REL).//0.66:23:52//HOMO SAPIENS (HUMAN).//P36269
 F-NT2RP2003706//GLUTAMYL AMINOPEPTIDASE (EC 3.4.11.7) (EAP) (AMINOPEPTIDASE A) (APA) (DIFFERENTIATION ANTIGEN GP160).//1.2e-22:187:35//HOMO SAPIENS (HUMAN).//Q07075
 F-NT2RP2003713//UBIQUITIN CARBOXYL-TERMINAL HYDROLASE 6 (EC 3.1.2.15) (UBIQUITIN THIOLESTERASE 6) (UBIQUITIN-SPECIFIC PROCESSING PROTEASE 6) (DEUBIQUITINATING ENZYME 6) (PROTO-ONCOGENE TRE-2).//2.7e-06:119:34//HOMO SAPIENS (HUMAN).//P35125
 F-NT2RP2003714//ZINC FINGER PROTEIN 91 (ZINC FINGER PROTEIN HTF10) (HPF7).//6.7e-27:68:75//HOMO SAPIENS (HUMAN).//Q05481
 F-NT2RP2003727//HYPOTHETICAL PROTEIN MC007 HOMOLOG.//0.64:110:30//MYCOPLASMA PNEUMONIAE.//P75105
 F-NT2RP2003737//UBIQUITIN-CONJUGATING ENZYME E2-17 KD 2 (EC 6.3.2.19) (UBIQUITIN-PROTEIN LIGASE) (UBIQUITIN CARRIER PROTEIN) (E2(17)KB 2).//1.2e-72:147:90//HOMO SAPIENS (HUMAN), MUS MUSCULUS (MOUSE), RATTUS NORVEGICUS (RAT), AND XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P51669
 F-NT2RP2003751//EXTRACELLULAR GLOBIN PRECURSOR.//0.67:68:30//PSEUDOTERRANOVA DECIPIENS (COD WORM).//P26914

3013

F-NT2RP2003760//COATOMER GAMMA SUBUNIT (GAMMA-COAT PROTEIN) (GAMMA-COP).//1.0e-98:235:82//BOS TAURUS (BOVINE).//P53620

F-NT2RP2003764//HYPOTHETICAL 29.3 KD PROTEIN (ORF9 2).//0.011:69:34//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPMNPV).//010341

F-NT2RP2003769//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3 4) (A6L).//1.0:33:36//EQUUS CABALLUS (HORSE).//P48663

F-NT2RP2003770//PHOSPHATE REGULON SENSOR PROTEIN P HOR (EC 2.7.3.-) (FRAGMENT).//0.029:35:42//PSEUDOMONAS AERUGINOSA.//P23621

F-NT2RP2003777//HYPOTHETICAL 82 KD AVIRULENCE PROTEIN IN AVRBS3 REGION.//0.041:67:34//XANTHOMONAS CAMPESTRIS (PV. VESICATORIA).//P14728

F-NT2RP2003781//HYPOTHETICAL 36.7 KD PROTEIN AH6.2 IN CHROMOSOME II.//4.7e-54:204:47//CAENORHABDITIS ELEGANS.//Q09201

F-NT2RP2003793//PSEUDO-HEVEIN (MINOR HEVEIN).//0.61:30:36//HEVEA BRASILIENSIS (PARA RUBBER TREE).//P80359

F-NT2RP2003825//ENDOTHELIN-1 PRECURSOR (ET-1) (FRAGMENT).//1.0:35:37//CANIS FAMILIARIS (DOG).//P13206

F-NT2RP2003840//HYPOTHETICAL 48.1 KD PROTEIN B040 3.2 IN CHROMOSOME X.//2.5e-05:80:38//CAENORHABDITIS ELEGANS.//Q11076

F-NT2RP2003857//BACTERIOCIN MICROCIN B17 PRECURSOR (MCB17).//0.54:28:50//ESCHERICHIA COLI.//P05834

F-NT2RP2003859//DROSOCIN PRECURSOR.//1.0:37:35//DROSOPHILA MELANOGASTER (FRUIT FLY).//P36193

F-NT2RP2003871

F-NT2RP2003885//CUTICLE PROTEIN 32 (LM-32) (LM-ACP 32) (FRAGMENT).//1.0:28:50//LOCUSTA MIGRATORIA (MIGRATORY LOCUST).//P11736

F-NT2RP2003912//SERINE/THREONINE-PROTEIN KINASE NEK1 (EC 2.7.1.-) (NIMA-RELATED PROTEIN KINASE 1).//4.8e-110:268:80//MUS MUSCULUS (MOUSE).//P51954

F-NT2RP2003952//AMINOPEPTIDASE B (EC 3.4.11.6) (ARGINYL AMINOPEPTIDASE) (ARGININE AMINOPEPTIDASE) (CYTOSOL AMINOPEPTIDASE IV) (AP-B).//0.00024:92:31//RATTUS NORVEGICUS (RAT).//009175

F-NT2RP2003968//SKIN SECRETORY PROTEIN XP2 PRECURSOR (APEG PROTEIN).//9.2e-05:101:36//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P17437

F-NT2RP2003976//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//1.7e-21:62:62//HOMO SAPIENS (HUMAN).//P39188

F-NT2RP2003981//VACUOLAR PROTEIN SORTING-ASSOCIATE D PROTEIN VPS8.//2.7e-08:165:22//SACCHAROMYCES CER

3014

EVISIAE (BAKER'S YEAST).//P39702

F-NT2RP2003984//UNC-87 PROTEIN.//0.75:71:28//CAENORHABDITIS ELEGANS.//P37806

F-NT2RP2003986//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//5.3e-19:47:70//HOMO SAPIENS (HUMAN).//P39193

F-NT2RP2003988//!!!! ALU SUBFAMILY SX WARNING ENTRY !!!!!//2.2e-18:80:58//HOMO SAPIENS (HUMAN).//P39195

F-NT2RP2004013//TRANSCRIPTION FACTOR BTF3 (RNA POLYMERASE B TRANSCRIPTION FACTOR 3).//1.0e-52:141:77//HOMO SAPIENS (HUMAN).//P20290

F-NT2RP2004014//MACROPHAGE INFLAMMATORY PROTEIN-2-ALPHA (MIP2-ALPHA) (CINC-2-ALPHA).//0.99:45:26//RATTUS NORVEGICUS (RAT).//Q10746

F-NT2RP2004041//SYNAPSINS IA AND IB.//0.0022:51:37//BOS TAURUS (BOVINE).//P17599

F-NT2RP2004042//CRUSTACEAN HYPERGLYCEMIC HORMONE P RECURSOR (CHH) (FRAGMENT).//1.0:49:28//PENAEUS VAN NAMEI (PENOEID SHRIMP) (EUROPEAN WHITE SHRIMP).//Q26181

F-NT2RP2004066//CALDESMON (CDM).//2.9e-05:175:21//GALLUS GALLUS (CHICKEN).//P12957

F-NT2RP2004081//CADMIUM-METALLOTHIONEIN (CD-MT).//0.93:59:23//HELIX POMATIA (ROMAN SNAIL) (EDIBLE SNAIL).//P33187

F-NT2RP2004098//RAS SUPPRESSOR PROTEIN 1 (RSU-1) (RSP-1 PROTEIN) (RSP-1).//4.6e-09:121:30//HOMO SAPIENS (HUMAN).//Q15404

F-NT2RP2004124//NONHISTONE CHROMOSOMAL PROTEIN HMG-17.//0.068:63:31//GALLUS GALLUS (CHICKEN).//P02314

F-NT2RP2004142//HYPOTHETICAL 59.1 KD PROTEIN IN VP S15-YMC2 INTERGENIC REGION.//7.9e-05:94:28//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38262

F-NT2RP2004152//LAMIN L(1).//0.25:167:19//XENOPUS LAEVIS (AFRICAN CLAWED FROG).//P09010

F-NT2RP2004165//HYPOTHETICAL 57.5 KD PROTEIN IN VM A7-RPS25A INTERGENIC REGION.//0.0014:124:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P53214

F-NT2RP2004170//PROLINE-RICH PROTEIN MP-3 (FRAGMENT).//0.012:125:30//MUS MUSCULUS (MOUSE).//P05143

F-NT2RP2004172//HYPOTHETICAL 105.7 KD PROTEIN IN TPK3-PIR1 INTERGENIC REGION.//4.1e-26:214:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P36051

F-NT2RP2004187//ZINC FINGER PROTEIN 174.//3.7e-12:76:47//HOMO SAPIENS (HUMAN).//Q15697

F-NT2RP2004194//HYPOTHETICAL 10.5 KD PROTEIN C31A 2.13C IN CHROMOSOME I.//0.0013:92:23//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q09730

3015

F-NT2RP2004196//METALLOTHIONEIN 10-II (MT-10-II).//
 /0.92:36:36//MYTILUS EDULIS (BLUE MUSSEL).//P80247
 F-NT2RP2004207//MALE ACCESSORY GLAND SECRETORY PRO
 TEIN 355A PRECURSOR.//0.92:62:35//DROSOPHILA SIMUL
 ANS (FRUIT FLY).//P33737
 F-NT2RP2004226//66 KD STRESS PROTEIN (P66).//0.03
 0:113:26//PHYSARUM POLYCEPHALUM (SLIME MOLD).//P90
 587
 F-NT2RP2004232//PROTEIN KINASE C, MU TYPE (EC 2.7.
 1.-) (NPKC-MU).//2.0e-48:211:51//HOMO SAPIENS (HUM 10
 AN).//Q15139
 F-NT2RP2004239//GLUTENIN, HIGH MOLECULAR WEIGHT SU
 BUNIT PW212 PRECURSOR.//0.00038:111:36//TRITICUM A
 ESTIVUM (WHEAT).//P08489
 F-NT2RP2004240//METALLOTHIONEIN-II (MT-II) (METALL
 OTHIONEIN-LIKE PROTEIN) (MT-CE).//1.0:39:28//CAENO
 RHABDITIS ELEGANS.//P17512
 F-NT2RP2004242//RAS-RELATED PROTEIN RGP1 (GTP-BIND
 ING REGULATORY PROTEINRGP1).//0.0036:64:28//ORYZA
 SATIVA (RICE).//P25766
 F-NT2RP2004245//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
 4) (A6L).//1.0:21:42//PONGO PYGMAEUS PYGMAEUS (BOR
 NEAN ORANGUTAN).//P92896
 F-NT2RP2004270//SPIDROIN 2 (DRAGLINE SILK FIBROIN
 2) (FRAGMENT).//0.00023:118:33//NEPHILA CLAVIPES
 (ORB SPIDER).//P46804
 F-NT2RP2004300//PROBABLE E4 PROTEIN.//0.18:77:40//
 HUMAN PAPILOMAVIRUS TYPE 8.//P06425
 F-NT2RP2004316
 F-NT2RP2004321//HYPOTHETICAL 10.8 KD PROTEIN SSR24 30
 39.//1.0:50:28//SYNECHOCYSTIS SP. (STRAIN PCC 680
 3).//Q01904
 F-NT2RP2004339//!!!! ALU SUBFAMILY SX WARNING ENTR
 Y !!!!!/5.0e-33:84:77//HOMO SAPIENS (HUMAN).//P391
 95
 F-NT2RP2004347//HYPOTHETICAL 40.9 KD PROTEIN F33H
 1.3 FROM CHROMOSOME II.//0.78:96:30//CAENORHABDITI
 S ELEGANS.//Q09556
 F-NT2RP2004364//MINOR OUTER CAPSID PROTEIN (NS26)
 (NONSTRUCTURAL PROTEINVP9).//0.059:143:30//BOVINE 40
 ROTAVIRUS (STRAIN UK).//P04515
 F-NT2RP2004365//EAMZP30-47 PROTEIN (FRAGMENT).//0.
 27:38:39//EIMERIA ACERVULINA.//P21959
 F-NT2RP2004366//GLYCOPROTEIN L PRECURSOR.//0.64:7
 1:28//MAREK'S DISEASE HERPESVIRUS (STRAIN GA) (MDH
 V).//P52510
 F-NT2RP2004373//HISTIDINE-RICH GLYCOPROTEIN PRECUR
 SOR (HISTIDINE-PROLINERICH GLYCOPROTEIN) (HPRC) (F
 RAGMENT).//0.59:50:40//ORYCTOLAGUS CUNICULUS (RABB
 IT).//Q28640

3016

F-NT2RP2004389//HYPOTHETICAL 70.7 KD PROTEIN F09G
 8.3 IN CHROMOSOME III.//4.0e-16:89:43//CAENORHABDI
 TIS ELEGANS.//P34388
 F-NT2RP2004392
 F-NT2RP2004396//SINGLE-STRANDED NUCLEIC ACID-BINDI
 NG PROTEIN.//0.42:89:29//SACCHAROMYCES CEREVISIAE
 (BAKER'S YEAST).//P10080
 F-NT2RP2004399//SOMATOTROPIN PRECURSOR (GROWTH HOR
 MONE).//1.0:72:34//MESOCRICETUS AURATUS (GOLDEN HA
 MSTER).//P37886
 F-NT2RP2004400
 F-NT2RP2004412//SPERM PROTAMINE P1.//0.24:38:31//N
 OTORYCTES TYPHLOPS (MARSUPIAL MOLE).//P42143
 F-NT2RP2004425//SUPPRESSOR PROTEIN SRP40.//0.0087:
 197:22//SACCHAROMYCESCEREVISIAE (BAKER'S YEAST).//
 P32583
 F-NT2RP2004463//ALPHA-2A ADRENERGIC RECEPTOR (ALPH
 A-2A ADRENOCEPTOR) (ALPHA-2AAR).//1.3e-05:121:37//
 MUS MUSCULUS (MOUSE).//Q01338
 20 F-NT2RP2004476//NICKEL-SENSITIVE T-TYPE CALCIUM CH
 ANNEL ALPHA-1 SUBUNIT(RBE-II).//0.20:68:36//RATTUS
 NORVEGICUS (RAT).//Q07652
 F-NT2RP2004490//FOS-RELATED ANTIGEN 1.//0.94:59:33
 //HOMO SAPIENS (HUMAN).//P15407
 F-NT2RP2004512//NADH-UBIQUINONE OXIDOREDUCTASE CHA
 IN 4 (EC 1.6.5.3) (FRAGMENTS).//1.0:37:32//PISASTE
 R OCHRACEUS (SEA STAR).//P24998
 F-NT2RP2004523//!!!! ALU SUBFAMILY SQ WARNING ENTR
 Y !!!!!/2.1e-15:57:71//HOMO SAPIENS (HUMAN).//P391
 94
 F-NT2RP2004538//KINESIN-LIKE PROTEIN KIF1A (AXONAL
 TRANSPORTER OF SYNAPTIC VESICLES).//1.2e-48:121:6
 0//HOMO SAPIENS (HUMAN).//Q12756
 F-NT2RP2004551//HYPOTHETICAL 7.6 KD PROTEIN (ORF 6
 5).//1.0:20:50//EUGLENA GRACILIS.//P32095
 F-NT2RP2004568//PUTATIVE ATP-DEPENDENT RNA HELICAS
 E C30D11.03.//5.2e-07:150:30//SCHIZOSACCHAROMYCES
 POMBE (FISSION YEAST).//Q09903
 F-NT2RP2004580//!!!! ALU SUBFAMILY SC WARNING ENTR
 Y !!!!!/3.7e-37:100:78//HOMO SAPIENS (HUMAN).//P39
 192
 F-NT2RP2004587//HYPOTHETICAL 57.5 KD PROTEIN IN VM
 A7-RPS25A INTERGENIC REGION.//8.2e-06:150:32//SACC
 HAROMYCES CEREVISIAE (BAKER'S YEAST).//P53214
 F-NT2RP2004594//HYPOTHETICAL 45.3 KD PROTEIN C09F
 5.7 IN CHROMOSOME II.//0.84:105:24//CAENORHABDITIS
 ELEGANS.//Q09458
 F-NT2RP2004600//MYRISTOYLATED ALANINE-RICH C-KINAS
 E SUBSTRATE (MARCKS).//0.17:127:29//RATTUS NORVEGI
 CUS (RAT).//P30009

3017

F-NT2RP2004602//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/1.1e-05:50:58//HOMO SAPIENS (HUMAN).//P3918
 8
 F-NT2RP2004614//HYPOTHETICAL 11.6 KD PROTEIN.//1.
 0:68:33//VACCINIA VIRUS(STRAIN COPENHAGEN).//P2056
 1
 F-NT2RP2004655//GLYCINE-RICH RNA-BINDING PROTEIN
 7.//7.0e-05:70:42//ARABIDOPSIS THALIANA (MOUSE-EAR
 CRESS).//Q03250
 F-NT2RP2004664//HYPOTHETICAL 104.0 KD PROTEIN C32A 10
 11.03C IN CHROMOSOME I.//0.30:78:38//SCHIZOSACCHAR
 OMYCES POMBE (FISSION YEAST).//Q10328
 F-NT2RP2004675
 F-NT2RP2004681
 F-NT2RP2004689//HYPOTHETICAL 78.3 KD PROTEIN IN RA
 M2-ATP7 INTERGENIC REGION.//0.021:179:24//SACCHARO
 MYCES CEREVISIAE (BAKER'S YEAST).//P34243
 F-NT2RP2004709//HYPOTHETICAL PROTEIN MJ0647.//0.9
 0:39:43//METHANOCOCCUSJANNASCHII.//Q58063
 F-NT2RP2004710//GAR2 PROTEIN.//0.085:60:30//SCHIZO 20
 SACCHAROMYCES POMBE (FISSION YEAST).//P41891
 F-NT2RP2004736//!!!! ALU SUBFAMILY J WARNING ENTRY
 !!!!!/4.4e-15:97:49//HOMO SAPIENS (HUMAN).//P3918
 8
 F-NT2RP2004743//MALE SPECIFIC SPERM PROTEIN MST87
 F.//0.43:24:41//DROSOPHILA MELANOGASTER (FRUIT FL
 Y).//P08175
 F-NT2RP2004767//36.4 KD PROLINE-RICH PROTEIN.//0.0
 051:88:27//LYCOPERSICON ESCULENTUM (TOMATO).//Q004
 51
 F-NT2RP2004768//SERINE/THREONINE-PROTEIN KINASE NR
 K1 (EC 2.7.1.-) (N-RICH KINASE 1).//9.0e-29:166:43
 //SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P3869
 2
 F-NT2RP2004775
 F-NT2RP2004791//PROBABLE LEUCYL-TRNA SYNTHETASE (E
 C 6.1.1.4) (LEUCINE--TRNA LIGASE) (LEURS).//7.4e-6
 0:226:53//CAENORHABDITIS ELEGANS.//Q09996
 F-NT2RP2004799//SUCCINYL-COA LIGASE [GDP-FORMING],
 BETA-CHAIN PRECURSOR(EC 6.2.1.4) (SUCCINYL-COA SY 40
 NTHETASE, BETA CHAIN) (SCS-BETA).//2.2e-42:133:57/
 /NEOCALLIMASTIX FRONTALIS (RUMEN FUNGUS).//P53587
 F-NT2RP2004802//HYPOTHETICAL 17.1 KD PROTEIN IN PU
 R5 3' REGION.//0.018:86:32//SACCHAROMYCES CEREVISIA
 E (BAKER'S YEAST).//P38898
 F-NT2RP2004816//H58 PROTEIN.//1.0e-68:145:93//MUS
 MUSCULUS (MOUSE).//P40336
 F-NT2RP2004841//DSRD PROTEIN.//0.83:33:39//ARCHAEO
 GLOBUS FULCIDUS.//P70742
 F-NT2RP2004861//KERATIN, HIGH-SULFUR MATRIX PROTEI 50

3018

N, IIIA3A.//0.0072:41:39//OVIS ARIES (SHEEP).//P02
 443
 F-NT2RP2004897//METALLOTHIONEIN-LIKE PROTEIN 1.//
 0.99:41:41//CASUARINA GLAUCA (SWAMP OAK).//Q39511
 F-NT2RP2004933//DEATH-ASSOCIATED PROTEIN KINASE 1
 (EC 2.7.1.-) (DAP KINASE 1).//8.4e-34:102:67//HOMO
 SAPIENS (HUMAN).//P53355
 F-NT2RP2004936//HIGH POTENTIAL IRON-SULFUR PROTEI
 N, ISOZYME 2 (HIPIP 2).//0.87:36:33//ECTOTHIORHODO
 SPIRA VACUOLATA.//P38524
 F-NT2RP2004959//STEM CELL FACTOR PRECURSOR (SCF)
 (MAST CELL GROWTH FACTOR) (MGF) (C-KIT LIGAND).//
 1.0:69:28//CANIS FAMILIARIS (DOG).//Q06220
 F-NT2RP2004961//ZINC FINGER PROTEIN 33A (ZINC FING
 ER PROTEIN KOX31) (KIAA0065) (HA0946) (FRAGMENT)./
 /2.1e-21:73:58//HOMO SAPIENS (HUMAN).//Q06730
 F-NT2RP2004962//!!!! ALU SUBFAMILY SB WARNING ENTR
 Y !!!!!/0.17:28:57//HOMO SAPIENS (HUMAN).//P39189
 F-NT2RP2004967//HYPOTHETICAL 7.3 KD PROTEIN.//0.7
 6:41:31//THERMOPROTEUSTENAX VIRUS 1 (STRAIN KRA1)
 (TTV1).//P19301
 F-NT2RP2004978//SPERMATID-SPECIFIC PROTEIN T2 [CON
 TAINS: SPERM PROTAMINESP2].//0.44:40:45//SEPIA OFF
 ICINALIS (COMMON CUTTLEFISH).//P80002
 F-NT2RP2004982
 F-NT2RP2004985//HYPOTHETICAL PROTEIN KIAA0144.//1.
 2e-51:204:57//HOMO SAPIENS (HUMAN).//Q14157
 F-NT2RP2004999//LONG NEUROTOXIN 1 (ALPHA-BUNGAROTO
 XIN) (BGTX).//0.23:73:26//BUNGARUS MULTICINCTUS (M
 ANY-BANDED KRAIT).//P01378
 F-NT2RP2005000//ATPASE STABILIZING FACTOR 15 KD PR
 OTEIN.//0.12:37:32//SACCHAROMYCES CEREVISIAE (BAKE
 R'S YEAST).//P16965
 F-NT2RP2005001//SMALL PROLINE RICH PROTEIN II (SPR
 -II) (CLONE 930).//0.90:54:31//HOMO SAPIENS (HUMA
 N).//P22531
 F-NT2RP2005003//DOWN REGULATORY PROTEIN OF INTERLE
 UKIN 2 RECEPTOR.//1.6e-30:78:56//MUS MUSCULUS (MOU
 SE).//P15533
 F-NT2RP2005012//NPL1 PROTEIN (SEC63 PROTEIN).//0.0
 0024:94:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAS
 T).//P14906
 F-NT2RP2005018//GAG POLYPROTEIN (CORE POLYPROTEIN)
 [CONTAINS: CORE PROTEINS P19, P10] (FRAGMENT).//
 1.0:91:28//AVIAN ENDOGENOUS ROUS-ASSOCIATED VIRUS-
 0 (EV-2) (AVIAN RETROVIRUS RAV-0).//P06937
 F-NT2RP2005020
 F-NT2RP2005022//VEGETATIBLE INCOMPATIBILITY PROTEI
 N HET-E-1.//4.9e-11:106:35//PODOSPORA ANSERINA.//Q
 00808

- F-NT2RP2005031
F-NT2RP2005037//ANTI-SILENCING PROTEIN 1.//2.2e-3
2:117:55//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).
//P32447
F-NT2RP2005038//DNA NUCLEOTIDYLEXOTRANSFERASE (EC
2.7.7.31) (TERMINAL ADDITION ENZYME) (TERMINAL DEO
XYNUCLEOTIDYLTRANSFERASE) (TERMINAL TRANSFERASE)./
/9.3e-28:187:40//AMBYSTOMA MEXICANUM (AXOLOTL).//0
57486
F-NT2RP2005108//CUTICLE COLLAGEN 2.//0.33:62:38//C 10
AENORHABDITIS ELEGANS.//P17656
F-NT2RP2005116//PUTATIVE EUKARYOTIC TRANSLATION IN
ITIATION FACTOR 3 ALPHA SUBUNIT (EIF-3 ALPHA).//4.
0e-54:161:63//CAENORHABDITIS ELEGANS.//P34466
F-NT2RP2005126//CHLOROPLAST 50S RIBOSOMAL PROTEIN
L27 (FRAGMENT).//0.23:46:39//PLEUROCHRYSID HAPTONE
MOFERA.//P41552
F-NT2RP2005139//COLLAGEN ALPHA 1(XI1) CHAIN (FRAGM
ENTS).//0.016:43:37//BOS TAURUS (BOVINE).//P25508
F-NT2RP2005140//HYPOTHETICAL 7.4 KD PROTEIN YCF33. 20
//0.96:51:39//GUILLARDIA THETA (CRYPTOMONAS PHI)./
/078517
F-NT2RP2005144//TUBBY PROTEIN.//5.6e-08:66:45//MUS
MUSCULUS (MOUSE).//P50586
F-NT2RP2005147
F-NT2RP2005159//PHOTOSYSTEM II 4 KD REACTION CENTR
E PROTEIN PRECURSOR.//0.94:57:29//NICOTIANA TABACU
M (COMMON TOBACCO), AND SPINACIA OLERACEA (SPINAC
H).//P12164
F-NT2RP2005162//HYPOTHETICAL 54.2 KD PROTEIN IN ER 30
P5-ORC6 INTERGENIC REGION.//1.2e-33:139:51//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P38821
F-NT2RP2005168//HETEROGENOUS NUCLEAR RIBONUCLEOPRO
TEIN U (HNRNP U).//2.8e-33:102:61//HOMO SAPIENS (H
UMAN).//Q00839
F-NT2RP2005204//DNA DAMAGE TOLERANCE PROTEIN RHC31
(RAD31 HOMOLOG).//3.9e-28:141:42//SACCHAROMYCES C
EREVISIAE (BAKER'S YEAST).//Q06624
F-NT2RP2005227
F-NT2RP2005239//TRNA SPLICING PROTEIN SPL1.//2.0e- 40
38:117:64//CANDIDA ALBICANS (YEAST).//P87185
F-NT2RP2005254//OMEGA-AGATOXIN IB (OMEGA-AGA-IB)
(FRAGMENT).//0.26:29:48//AGELLENOPSIS APERTA (FUNNE
L-WEB SPIDER).//P15970
F-NT2RP2005270//HOMEBOX PROTEIN HOX-A4 (CHOX-1.
4).//0.037:82:34//GALLUSGALLUS (CHICKEN).//P17277
F-NT2RP2005276//LONG-CHAIN-FATTY-ACID--COA LIGASE
4 (EC 6.2.1.3) (LONG-CHAIN ACYL-COA SYNTHETASE 4)
(LACS 4).//2.0e-59:174:61//RATTUS NORVEGICUS(RAT).
//035547
F-NT2RP2005287//ZINC FINGER PROTEIN 26 (ZINC FINGE
R PROTEIN KOX20) (FRAGMENT).//1.5e-05:27:70//HOMO
SAPIENS (HUMAN).//P17031
F-NT2RP2005288//PROBABLE RUBREDOXIN HUPI.//1.0:42:
28//RHIZOBIUM LEGUMINOSARUM (BIOVAR VICIAE).//P281
51
F-NT2RP2005289//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!//1.1e-21:75:70//HOMO SAPIENS (HUMAN).//P391
93
F-NT2RP2005293//TRANSLATION INITIATION FACTOR IF-
2.//0.58:170:24//HELICOBACTER PYLORI (CAMPYLOBACTE
R PYLORI).//P55972
F-NT2RP2005315//CUTICLE COLLAGEN 7 (FRAGMENT).//0.
091:65:38//CAENORHABDITIS ELEGANS.//P18832
F-NT2RP2005325//CHROMOGRANIN A PRECURSOR (CGA) (PI
TUITARY SECRETORY PROTEIN I) (SP-I) [CONTAINS: PAN
CREASTATIN; WE-14].//9.5e-09:98:39//HOMO SAPIENS
(HUMAN).//P10645
F-NT2RP2005336//HYPOTHETICAL 68.7 KD PROTEIN IN ST
B1-MCK1 INTERGENIC REGION.//0.00011:124:28//SACCHA
ROMYCES CEREVISIAE (BAKER'S YEAST).//P42846
F-NT2RP2005344//PROBABLE CALCIUM-TRANSPORTING ATPA
SE 4 (EC 3.6.1.38).//4.7e-21:92:52//SACCHAROMYCES
CEREVISIAE (BAKER'S YEAST).//Q12675
F-NT2RP2005354
F-NT2RP2005358//MYOSIN IC HEAVY CHAIN.//0.012:91:3
9//ACANTHAMOEBA CASTELLANII (AMOEBA).//P10569
F-NT2RP2005360//ACROSIN PRECURSOR (EC 3.4.21.10)./
/0.0022:73:36//ORYCTOLAGUS CUNICULUS (RABBIT).//P4
8038
F-NT2RP2005393//HYPOTHETICAL 25.9 KD PROTEIN AH6.3
IN CHROMOSOME II.//0.00085:135:28//CAENORHABDITIS
ELEGANS.//Q09202
F-NT2RP2005407//SQUALENE MONOOXYGENASE (EC 1.14.9
9.7) (SQUALENE EPOXIDASE) (SE).//0.96:109:26//SACC
HAROMYCES CEREVISIAE (BAKER'S YEAST).//P32476
F-NT2RP2005436//EXTENSIN PRECURSOR (PROLINE-RICH G
LYCOPROTEIN).//0.0011:54:42//ZEA MAYS (MAIZE).//P1
4918
F-NT2RP2005441//PROLINE-RICH PROTEIN MP-2 PRECURSO
R.//0.039:182:29//MUSMUSCULUS (MOUSE).//P05142
F-NT2RP2005453
F-NT2RP2005457//NADH-UBIQUINONE OXIDOREDUCTASE SUB
UNIT B14.5B (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-
B14.5B) (CI-B14.5B).//4.0e-10:124:37//BOS TAURUS
(BOVINE).//Q02827
F-NT2RP2005464//HYPOTHETICAL 9.5 KD PROTEIN.//0.9
6:42:33//VACCINIA VIRUS(STRAIN COPENHAGEN).//P2055
3
50 F-NT2RP2005465//MITOCHONDRIAL CARRIER PROTEIN RIM

2. //4.6e-09:92:42//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P38127
 F-NT2RP2005472//HYPOTHETICAL PROTEIN BB0129. //0.76:80:32//BORRELIA BURGDORFERI (LYME DISEASE SPIROCHETE). //051155
 F-NT2RP2005476//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//1.0e-31:39:89//HOMO SAPIENS (HUMAN). //P39193
 F-NT2RP2005490//METALLOTHIONEIN-II (MT-II). //0.14:27:33//SCYLLA SERRATA (MUD CRAB). //P02806
 F-NT2RP2005491//DNA-DIRECTED RNA POLYMERASE SUBUNIT I (EC 2.7.7.6). //0.95:45:31//METHANOCOCCUS JANNASCHII. //Q58785
 F-NT2RP2005495//HYPOTHETICAL 10.8 KD PROTEIN IN CP30-R111 INTERGENIC REGION. //0.99:68:30//BACTERIOPHAGE T4. //Q02407
 F-NT2RP2005496//ZINC FINGER PROTEIN 135. //1.4e-54:120:59//HOMO SAPIENS (HUMAN). //P52742
 F-NT2RP2005498//PROTEIN PHOSPHATASE PP2A, 55 KD REGULATORY SUBUNIT, ALPHA ISOFORM (PROTEIN PHOSPHATASE PP2A B SUBUNIT ALPHA ISOFORM) (ALPHA-PR55). //9.5e-76:146:86//RATTUS NORVEGICUS (RAT). //P36876
 F-NT2RP2005501//GALECTIN-3 (GALACTOSE-SPECIFIC LECTIN 3) (MAC-2 ANTIGEN) (ICE-BINDING PROTEIN) (35 KD LECTIN) (CARBOHYDRATE BINDING PROTEIN 35) (CBP 35) (LAMININ-BINDING PROTEIN) (LECTIN L-29) (L-31) (GALACTOSIDE-BINDING PROTEIN) (GALBP). //0.025:70:40//HOMO SAPIENS (HUMAN). //P17931
 F-NT2RP2005509//PROCOLLAGEN ALPHA 1(I) CHAIN PRECURSOR. //1.0:166:27//GALLUS GALLUS (CHICKEN). //P02457
 F-NT2RP2005520//CHROMOSOME ASSEMBLY PROTEIN XCAP-E. //7.9e-45:118:79//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P50533
 F-NT2RP2005525//50S RIBOSOMAL PROTEIN L11. //1.0:47:27//BORRELIA BURGDORFERI (LYME DISEASE SPIROCHETE). //051354
 F-NT2RP2005531//PROTEIN-TYROSINE PHOSPHATASE MEG1 (EC 3.1.3.48) (PTPASE-MEG1) (MEG). //9.8e-13:84:45//HOMO SAPIENS (HUMAN). //P29074
 F-NT2RP2005539//RING CANAL PROTEIN (KELCH PROTEIN). //4.9e-10:90:33//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q04652
 F-NT2RP2005540//NUCLEOTIDE BINDING PROTEIN EXPZ. //0.36:119:21//BACILLUS SUBTILIS. //P39115
 F-NT2RP2005549//HYPOTHETICAL 32.0 KD PROTEIN C16C1 0.10 IN CHROMOSOME III. //6.0e-39:179:46//CAENORHABDITIS ELEGANS. //Q09253
 F-NT2RP2005555
 F-NT2RP2005557//HYPOTHETICAL 23.7 KD PROTEIN C13C 50
- 6.14 IN CHROMOSOME I. //4.9e-06:90:35//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09790
 F-NT2RP2005581
 F-NT2RP2005600//BASIC PROLINE-RICH PEPTIDE P-E (IB-9). //0.014:37:40//HOMO SAPIENS (HUMAN). //P02811
 F-NT2RP2005605//GONADOLIBERIN I PRECURSOR (LHRH I) (LUTEINIZING HORMONE-RELEASING HORMONE I) (GONADOTROPIN-RELEASING HORMONE I) (GNRH I) (LULIBERIN I) (FRAGMENT). //0.64:26:42//MACACA MULATTA (RHESUS MACAQUE). //P55247
 F-NT2RP2005620//HYPOTHETICAL 45.1 KD PROTEIN IN RP55-ZMS1 INTERGENIC REGION. //8.7e-31:138:49//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P47160
 F-NT2RP2005622//NEUROTOXIN-LIKE PROTEIN STR1 (ANATOXIN AAH STR1). //0.39:22:40//ANDROCTONUS AUSTRALIS HECTOR (SAHARA SCORPION). //P80950
 F-NT2RP2005635//HYPOTHETICAL 80.7 KD PROTEIN IN ERG7-NMD2 INTERGENIC REGION. //5.8e-43:144:56//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P38795
 F-NT2RP2005637//VPU PROTEIN (U ORF PROTEIN). //0.91:33:45//CHIMPANZEE IMMUNODEFICIENCY VIRUS (SIV(CPZ)) (CIV). //P17286
 F-NT2RP2005640//METALLOTHIONEIN-LIKE PROTEIN LSC5 4. //0.63:41:31//BRASSICA NAPUS (RAPE). //P43402
 F-NT2RP2005645
 F-NT2RP2005651//OCTAMER-BINDING TRANSCRIPTION FACTOR 3A (OCT-3A) (OCT-4). //0.0023:50:42//HOMO SAPIENS (HUMAN). //Q01860
 F-NT2RP2005654//HYPOTHETICAL 48.6 KD PROTEIN IN BE T1-PAN1 INTERGENIC REGION. //6.1e-16:76:44//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40564
 F-NT2RP2005669//METALLOTHIONEIN-II (MT-II). //0.76:16:50//SCYLLA SERRATA (MUD CRAB). //P02806
 F-NT2RP2005675//PUTATIVE ORAL CANCER SUPPRESSOR (DELETED IN ORAL CANCER-1). //6.5e-26:116:54//MESOCRICETUS AURATUS (GOLDEN HAMSTER). //P49119
 F-NT2RP2005683//HYPOTHETICAL PROTEIN HI0275. //0.17:50:40//HAEMOPHILUS INFLUENZAE. //P43975
 F-NT2RP2005690//PYRROLINE-5-CARBOXYLATE REDUCTASE (EC 1.5.1.2) (P5CR) (P5C REDUCTASE). //1.3e-16:75:30//PISUM SATIVUM (GARDEN PEA). //Q04708
 F-NT2RP2005694//HYPOTHETICAL PROTEIN KIAA0032. //9.6e-11:135:34//HOMO SAPIENS (HUMAN). //Q15034
 F-NT2RP2005701//SALIVARY PROLINE-RICH PROTEIN P0 (ALLELE M) [CONTAINS: PEPTIDE P-D] (FRAGMENT). //0.084:158:32//HOMO SAPIENS (HUMAN). //P10161
 F-NT2RP2005712//METALLOTHIONEIN-II (MT-II). //0.19:14:50//STENELLA COERULEOALBA (STRIPED DOLPHIN). //P14425
 F-NT2RP2005719//COLLAGEN ALPHA 1(I) CHAIN (FRAGMENT)

3023

T. //1.0:36:41//ORYCTOLAGUS CUNICULUS (RABBIT). //P02456
 F-NT2RP2005722//ZINC FINGER PROTEIN ZFP-36 (FRAGMENT). //7.8e-37:131:62//HOMO SAPIENS (HUMAN). //P16415
 F-NT2RP2005723//IIII ALU SUBFAMILY SC WARNING ENTRY IIIII//0.98:23:60//HOMO SAPIENS (HUMAN). //P39192
 F-NT2RP2005726//HYPOTHETICAL PROTEIN TP0375. //0.98:30:43//TREPONEMA PALLIDUM. //083390
 F-NT2RP2005732//PERIOD CLOCK PROTEIN (FRAGMENT). //100.41:20:55//DROSOPHILAROBUSTA (FRUIT FLY). //Q03296
 F-NT2RP2005741//SMR1 PROTEIN PRECURSOR (VCS-ALPHA 1). //0.38:58:36//RATTUS NORVEGICUS (RAT). //P13432
 F-NT2RP2005748//ZINC FINGER PROTEIN KOX23 (FRAGMENT). //0.026:19:68//HOMOSAPIENS (HUMAN). //P17034
 F-NT2RP2005752//PROCOLLAGEN ALPHA 1(III) CHAIN PRECURSOR. //0.90:101:31//HOMO SAPIENS (HUMAN). //P02461
 F-NT2RP2005753//DNA-BINDING P52/P100 COMPLEX, 100 KD SUBUNIT (FRAGMENTS). //0.50:22:59//HOMO SAPIENS (HUMAN). //P30808
 F-NT2RP2005763//PUTATIVE ATP-DEPENDENT RNA HELICASE STE13. //4.7e-14:108:37//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q09181
 F-NT2RP2005767//NONHISTONE CHROMOSOMAL PROTEIN 6B. //4.1e-08:65:32//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P11633
 F-NT2RP2005773//PYRROLINE-5-CARBOXYLATE REDUCTASE (EC 1.5.1.2) (P5CR) (P5C REDUCTASE). //1.2e-14:65:61//HOMO SAPIENS (HUMAN). //P32322
 F-NT2RP2005775//NEUROLYSIN PRECURSOR (EC 3.4.24.16) (NEUROTENSIN ENDOPEPTIDASE) (MITOCHONDRIAL OLIGOPEPTIDASE M) (MICROSOMAL ENDOPEPTIDASE) (MEP). //1.3e-103:199:90//ORYCTOLAGUS CUNICULUS (RABBIT). //P42675
 F-NT2RP2005781//SALIVARY ACIDIC PROLINE-RICH PHOSPHOPROTEIN 1/2 PRECURSOR (PRP-1 / PRP-3) (PRP-2 / PRP-4) (PIF-F / PIF-S) (PROTEIN A / PROTEIN C) [CONTAINS: PEPTIDE P-C]. //0.090:73:36//HOMO SAPIENS (HUMAN). //P02810
 F-NT2RP2005784//TRANS-ACTING TRANSCRIPTIONAL PROTEIN ICPO (IMMEDIATE-EARLY PROTEIN IE110) (VMW110) (ALPHA-0 PROTEIN). //3.5e-06:79:37//HERPES SIMPLEX VIRUS (TYPE 1 / STRAIN 17). //P08393
 F-NT2RP2005804//HYPOTHETICAL PROLINE-RICH PROTEIN (FRAGMENT). //1.8e-07:43:55//OWENIA FUSIFORMIS. //P21260
 F-NT2RP2005812//HYPOTHETICAL 39.3 KD PROTEIN IN CCN4-WBP1 INTERGENIC REGION. //6.3e-14:143:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40004

3024

F-NT2RP2005815//FERROCHELATASE (EC 4.99.1.1) (PROTOME FERRO-LYASE) (HEME SYNTHETASE). //0.0017:123:37//MYCOBACTERIUM AVIUM. //007401
 F-NT2RP2005835//SHP1 PROTEIN. //1.2e-08:135:26//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P34223
 F-NT2RP2005841//SMALL PROLINE-RICH PROTEIN II (SPR-II) (CLONE 174N). //0.23:28:53//HOMO SAPIENS (HUMAN). //P22532
 F-NT2RP2005853//HYPOTHETICAL 8.5 KD PROTEIN IN ASI A-MOTA INTERGENIC REGION. //0.99:33:48//BACTERIOPHAGE T4. //P22917
 F-NT2RP2005857//CHROMOSOME ASSEMBLY PROTEIN XCAP-C. //8.6e-84:235:66//XENOPUS LAEVIS (AFRICAN CLAWED FROG). //P50532
 F-NT2RP2005859//MALE SPECIFIC SPERM PROTEIN MST84D B. //0.017:60:40//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q01643
 F-NT2RP2005868//ATP SYNTHASE B' CHAIN PRECURSOR (EC 3.6.1.34) (SUBUNIT II). //0.28:121:28//SPINACIA OLERACEA (SPINACH). //P31853
 F-NT2RP2005886//MICRONUCLEAR LINKER HISTONE POLYPEPTIDE (MIC LH) [CONTAINS: LINKER HISTONE PROTEINS ALPHA, BETA, DELTA AND GAMMA]. //0.80:130:28//TETRAHYMENA THERMOPHILA. //P40631
 F-NT2RP2005890
 F-NT2RP2005901//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (A6L). //0.35:18:44//DROSOPHILA YAKUBA (FRUIT FLY). //P03933
 F-NT2RP2005908//IIII ALU SUBFAMILY SQ WARNING ENTRY IIIII//1.0e-28:61:65//HOMO SAPIENS (HUMAN). //P39194
 F-NT2RP2005933//PERIOD CLOCK PROTEIN (P230) (FRAGMENT). //1.7e-11:85:49//ACETABULARIA MEDITERRANEA (MERMAID'S WINE GLASS). //P12347
 F-NT2RP2005942//POLY(A) POLYMERASE (EC 2.7.7.19) (PAP) (POLYNUCLEOTIDE ADENYLYLTRANSFERASE). //7.2e-59:216:58//BOS TAURUS (BOVINE). //P25500
 F-NT2RP2005980//HYPOTHETICAL 11.5 KD PROTEIN IN RS P8A-AST1 INTERGENIC REGION. //1.0:49:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P38185
 F-NT2RP2006023//DNA REPAIR PROTEIN RECN (RECOMBINATION PROTEIN N) (FRAGMENT). //1.0:40:45//VIBRIO CHOLERAE. //P52118
 F-NT2RP2006038//HYPOTHETICAL 30.2 KD PROTEIN C02F 5.4 IN CHROMOSOME III. //4.0e-11:90:34//CAENORHABDITIS ELEGANS. //P34281
 F-NT2RP2006043//LAMININ BETA-1 CHAIN VARIANT (LAMININ BETA-1-2 CHAIN) (FRAGMENT). //0.00067:73:38//GALLUS GALLUS (CHICKEN). //Q01636
 F-NT2RP2006052//METALLOTHIONEIN-I (MT-I). //0.19:3

- 1:38//CERCOPITHECUS AETHIOPS (GREEN MONKEY) (GRIVE T).//P02797
 F-NT2RP2006069//COLLAGEN ALPHA 2(I) CHAIN (FRAGMEN TS).//1.0:66:34//RATTUS NORVEGICUS (RAT).//P02466
 F-NT2RP2006071//RESTIN.//0.40:156:29//GALLUS GALLUS (CHICKEN).//042184
 F-NT2RP2006098//HYPOTHETICAL 21.7 KD PROTEIN IN TU P1-ABP1 INTERGENIC REGION.//0.99:95:20//SACCHAROMY CES CEREVISIAE (BAKER'S YEAST).//P25651
 F-NT2RP2006100//LONG NEUROTOXIN 4 (ALPHA-NEUROTOXI N).//0.94:43:34//OPHIOPHAGUS HANNAH (KING COBRA) (NAJA HANNAH).//P80156
 F-NT2RP2006103//50S RIBOSOMAL PROTEIN L32.//0.40:3 6:38//SYNECHOCYSTIS SP. (STRAIN PCC 6803).//P73014
 F-NT2RP2006106//CUTICLE COLLAGEN 1.//0.28:85:29//C AENORHABDITIS ELEGANS.//P08124
 F-NT2RP2006141//HYPOTHETICAL 72.5 KD PROTEIN C2F7. 10 IN CHROMOSOME I.//1.9e-08:57:42//SCHIZOSACCHARO MYCES POMBE (FISSION YEAST).//Q09701
 F-NT2RP2006166
 F-NT2RP2006184//HYPOTHETICAL 11.2 KD PROTEIN IN CS GC-MDOG INTERGENIC REGION PRECURSOR.//0.95:87:26// ESCHERICHIA COLI.//P75917
 F-NT2RP2006186//MICROTUBULE-ASSOCIATED PROTEIN 2. /0.088:124:33//MUS MUSCULUS (MOUSE).//P20357
 F-NT2RP2006196//!!!! ALU SUBFAMILY SP WARNING ENTR Y !!!!!/4.0e-05:49:61//HOMO SAPIENS (HUMAN).//P391 93
 F-NT2RP2006200//PROCOLLAGEN ALPHA 2(V) CHAIN PRECU RSOR.//0.0013:205:32//HOMO SAPIENS (HUMAN).//P0599 7
 F-NT2RP2006219//GONADAL PROTEIN GDL.//3.5e-18:158: 37//DROSOPHILA MELANOGASTER (FRUIT FLY).//P22468
 F-NT2RP2006237//FIBRINOGEN- AND IG-BINDING PROTEIN PRECURSOR (MRP PROTEIN).//0.79:103:28//STREPTOCO CUS PYOGENES.//P30141
 F-NT2RP2006238//PROLINE-RICH PROTEIN MP-3 (FRAGMEN T).//4.7e-07:127:39//MUS MUSCULUS (MOUSE).//P05143
 F-NT2RP2006258//PROBABLE E5 PROTEIN.//0.78:47:34// RHESUS PAPILLOMAVIRUSTYPE 1 (RHPV 1).//P24834
 F-NT2RP2006261//PENAEIDIN-3A PRECURSOR (P3-A).//0. 61:35:40//PENAEUS VANNAMEI (PENOEID SHRIMP) (EUROP EAN WHITE SHRIMP).//P81058
 F-NT2RP2006275//ELECTROMOTOR NEURON-ASSOCIATED PRO TEIN 2 (FRAGMENT).//1.2e-28:59:57//TORPEDO CALIFOR NICA (PACIFIC ELECTRIC RAY).//P14401
 F-NT2RP2006312//HIGH-MOBILITY-GROUP PROTEIN (NONHI STONE CHROMOSOMAL PROTEIN).//1.6e-06:53:35//TETRAH YMENA PYRIFORMIS.//P40625
 F-NT2RP2006320//TAT PROTEIN (TRANSACTIVATING REGUL ATORY PROTEIN) (FRAGMENT).//0.90:24:41//HUMAN IMMU NODEFICIENCY VIRUS TYPE 1 (BH5 ISOLATE) (HIV-1).// P04612
 F-NT2RP2006321//!!!! ALU SUBFAMILY SP WARNING ENTR Y !!!!!/0.0051:25:76//HOMO SAPIENS (HUMAN).//P3919 3
 F-NT2RP2006323//WISKOTT-ALDRICH SYNDROME PROTEIN (WASP).//0.84:33:39//HOMO SAPIENS (HUMAN).//P42768
 F-NT2RP2006333//MYOTOXIN 3 PRECURSOR (CROTAMINE 3).//0.56:37:40//CROTALUS DURISSUS TERRIFICUS (SOU TH AMERICAN RATTLESNAKE).//P24333
 F-NT2RP2006334//SUCCINYL-COA LIGASE [GDP-FORMING], ALPHA-CHAIN 3 PRECURSOR (EC 6.2.1.4) (SUCCINYL-CO A SYNTHETASE, ALPHA CHAIN 3).//0.00097:46:41//TRIC HOMONAS VAGINALIS.//P53401
 F-NT2RP2006365//NONSPECIFIC LIPID-TRANSFER PROTEIN 4.3 PRECURSOR (LTP 4.3).//0.18:75:29//HORDEUM VUL GARE (BARLEY).//Q42842
 F-NT2RP2006393//OMEGA-CONOTOXIN MVIIC PRECURSOR (F RAGMENT).//0.82:15:66//CONUS MAGUS (MAGUS CONE).// P37300
 F-NT2RP2006436//ANTERIOR-RESTRICTED HOMEBOX PROTE IN (RATHEKE POUCH HOMEBOX).//1.4e-08:50:50//MUS MU SCULUS (MOUSE).//Q61658
 F-NT2RP2006441//METALLOTHIONEIN-LIKE PROTEIN 1.// 0.99:22:54//MIMULUS GUTTATUS (SPOTTED MONKEY FLOWE R) (YELLOW MONKEY FLOWER).//P20238
 F-NT2RP2006454//SPERM PROTAMINE P1.//0.60:47:36//T ACHYGLOSSUS ACULEATUSACULEATUS (AUSTRALIAN ECHIDN A).//P35311
 F-NT2RP2006456
 F-NT2RP2006464//PHOTOSYSTEM I IRON-SULFUR CENTER (PHOTOSYSTEM I SUBUNITVII) (9 KD POLYPEPTIDE) (PSI -C).//0.91:79:30//SYNECHOCOCCUS SP. (STRAINPCC 700 2) (ACMENELLUM QUADRUPLICATUM).//P31087
 F-NT2RP2006467//PUTATIVE CUTICLE COLLAGEN F55C10. 3.//0.15:53:35//CAENORHABDITIS ELEGANS.//Q21184
 F-NT2RP2006472//HYPOTHETICAL 19 KD PROTEIN (ORF 16 7).//0.33:98:26//MARCHANTIA POLYMORPHA (LIVERWOR T).//P12202
 F-NT2RP2006534
 F-NT2RP2006554//ANTI-SIGMA F FACTOR ANTAGONIST (ST AGE II SPORULATION PROTEIN AA).//0.91:50:34//BACIL LUS SPHAERICUS.//032723
 F-NT2RP2006565//SECRETORY CARRIER-ASSOCIATED MEMBR ANE PROTEIN 1 (SCAMP 37).//6.0e-66:93:96//RATTUS N ORVEGICUS (RAT).//P56603
 F-NT2RP2006571//CYTOCHROME P450 2B10 (EC 1.14.14. 1) (CYP11B10) (TESTOSTERONE 16-ALPHA HYDROXYLASE) (P450-16-ALPHA) (CLONE PF3/46).//4.5e-40:138:57//M

US MUSCULUS (MOUSE). //P12791
 F-NT2RP2006573//SPERM PROTAMINE P1 (CYSTEINE-RICH PROTAMINE). //0.53:46:39//BOS TAURUS (BOVINE). //P02318
 F-NT2RP2006598//!!!! ALU SUBFAMILY SX WARNING ENTR Y !!!!!//1.3e-12:44:77//HOMO SAPIENS (HUMAN). //P39195
 F-NT2RP3000002//!!!! ALU SUBFAMILY SC WARNING ENTR Y !!!!!//1.4e-19:60:63//HOMO SAPIENS (HUMAN). //P39192
 F-NT2RP3000031//HYPOTHETICAL 89.8 KD PROTEIN F41H1 0.6 IN CHROMOSOME IV. //2.1e-39:210:42//CAENORHABDITIS ELEGANS. //Q20296
 F-NT2RP3000046//POSSIBLE THIOPHENE AND FURAN OXIDATION PROTEIN THDF. //1.4e-25:149:44//PSEUDOMONAS PUTIDA. //P25755
 F-NT2RP3000047//NPL4 PROTEIN. //4.7e-48:275:38//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P33755
 F-NT2RP3000050//ZINC FINGER PROTEIN 83 (ZINC FINGER PROTEIN HPF1). //3.2e-72:232:59//HOMO SAPIENS (HUMAN). //P51522
 F-NT2RP3000055//MALE SPECIFIC SPERM PROTEIN MST84DB. //0.26:57:36//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q01643
 F-NT2RP3000068//HYPOTHETICAL 182.0 KD PROTEIN IN NMD5-HOM6 INTERGENIC REGION. //0.0014:66:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P47170
 F-NT2RP3000072//HYPOTHETICAL 6.7 KD PROTEIN IN NOH A-CSPI INTERGENIC REGION. //0.95:49:30//ESCHERICHIA COLI. //P77695
 F-NT2RP3000080//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//5.1e-17:64:68//HOMO SAPIENS (HUMAN). //P39188
 F-NT2RP3000085//BIOTIN CARBOXYLASE (EC 6.3.4.14) (A SUBUNIT OF ACETYL-COA CARBOXYLASE (EC 6.4.1.2)) (ACC). //4.4e-43:169:51//BACILLUS SUBTILIS. //P49787
 F-NT2RP3000092//CELL DIVISION CONTROL PROTEIN 1. //0.00016:103:31//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40986
 F-NT2RP3000109//ACYL CARRIER PROTEIN HOMOLOG (ACP). //0.76:83:28//MYCOPLASMA GENITALIUM. //P47529
 F-NT2RP3000134
 F-NT2RP3000142//GAR2 PROTEIN. //0.00098:241:20//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //P41891
 F-NT2RP3000149//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (A6L). //0.0014:33:36//PONGO PYGMAEUS ABELII (SUMATRAN ORANGUTAN). //P92694
 F-NT2RP3000186//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//8.3e-15:36:83//HOMO SAPIENS (HUMAN). //P39188

8

F-NT2RP3000197//HYPOTHETICAL 6.0 KD PROTEIN IN THI 12 5' REGION. //0.91:21:52//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53820
 F-NT2RP3000207//GLUCOAMYLASE S1/S2 PRECURSOR (EC 3.2.1.3) (GLUCAN 1,4-ALPHA-GLUCOSIDASE) (1,4-ALPHA-D-GLUCAN GLUCOHYDROLASE). //0.026:209:27//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P08640
 F-NT2RP3000220//DNA-BINDING P52/P100 COMPLEX, 100 KD SUBUNIT (FRAGMENTS). //1.0:26:42//HOMO SAPIENS (HUMAN). //P30808
 F-NT2RP3000233//RING CANAL PROTEIN (KELCH PROTEIN). //2.1e-42:249:39//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q04652
 F-NT2RP3000235//HOMEBOX PROTEIN H40 (FRAGMENT). //0.55:45:40//APIS MELLIFERA (HONEYBEE). //P15858
 F-NT2RP3000247//HYPOTHETICAL PROTEIN KIA0218. //1.7e-82:123:69//HOMO SAPIENS (HUMAN). //Q93075
 F-NT2RP3000251//SERINE PROTEINASE STUBBLE (EC 3.4.21.-) (STUBBLE-STUBBLOID PROTEIN). //1.0:53:33//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q05319
 F-NT2RP3000252//HYPOTHETICAL 40 KD GTP-BINDING PROTEIN IN RIBOSOMAL PROTEIN GENE CLUSTER 5' REGION. //2.2e-06:96:32//HALOBACTERIUM CUTIRUBRUM. //P17103
 F-NT2RP3000255//HISTONE H1.1 (FRAGMENT). //0.95:71:33//BOS TAURUS (BOVINE). //P02253
 F-NT2RP3000267//HYPOTHETICAL 21.1 KD PROTEIN IN SS R-SERA INTERGENIC REGION (0182). //0.38:77:33//ESCHERICHIA COLI. //P09160
 F-NT2RP3000299//MYOSIN IC HEAVY CHAIN. //1.2e-11:147:34//ACANTHAMOEBA CASTELLANII (AMOEBA). //P10569
 F-NT2RP3000312//HYPOTHETICAL 57.5 KD PROTEIN IN VM A7-RPS25A INTERGENIC REGION. //0.64:216:29//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P53214
 F-NT2RP3000320//TRANSLATION INITIATION FACTOR IF-2. //5.2e-05:184:22//AQUIFEX AEOLICUS. //067825
 F-NT2RP3000324//HYPOTHETICAL PROTEIN HI1036. //0.69:64:35//HAEMOPHILUS INFLUENZAE. //P44097
 F-NT2RP3000333//WIRIA PROTEIN. //0.35:51:41//TRITICUM AESTIVUM (WHEAT). //Q01482
 F-NT2RP3000341//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//2.1e-30:57:80//HOMO SAPIENS (HUMAN). //P39189
 F-NT2RP3000348
 F-NT2RP3000350//HYPOTHETICAL 40 KD GTP-BINDING PROTEIN IN RIBOSOMAL PROTEIN GENE CLUSTER 5' REGION. //0.0011:77:35//HALOBACTERIUM CUTIRUBRUM. //P17103
 F-NT2RP3000359//GTP:AMP PHOSPHOTRANSFERASE MITOCHONDRIAL (EC 2.7.4.10) (AK3). //1.2e-97:222:84//BOS TAURUS (BOVINE). //P08760

3029

F-NT2RP3000361//PRE-MRNA SPLICING FACTOR PRP6.//2.
2e-08:128:28//SACCHAROMYCES CEREVISIAE (BAKER'S YE
AST).//P19735
F-NT2RP3000366//RAS-RELATED PROTEIN RAB-18.//2.1e-
107:206:99//MUS MUSCULUS (MOUSE).//P35293
F-NT2RP3000393//HOMEBOX PROTEIN HOX-C4 (HOX-3E)
(CP19).//0.0023:36:52//HOMO SAPIENS (HUMAN).//P090
17
F-NT2RP3000397//PUTATIVE PRE-MRNA SPLICING FACTOR
RNA HELICASE (DEAH BOXPROTEIN 13).//5.5e-27:116:44 10
//MUS MUSCULUS (MOUSE).//035286
F-NT2RP3000403//PRE-MRNA PROCESSING PROTEIN PRP40.
//0.00044:67:34//SACCHAROMYCES CEREVISIAE (BAKER'S
YEAST).//P33203
F-NT2RP3000418//RETROVIRUS-RELATED POLYPROTEIN
[CONTAINS: REVERSE TRANSCRIPTASE (EC 2.7.7.49); E
NDONUCLEASE].//2.2e-16:228:34//MUS MUSCULUS(MOUS
E).//P11369
F-NT2RP3000433//!!!! ALU SUBFAMILY J WARNING ENTRY
!!!!//1.7e-17:79:55//HOMO SAPIENS (HUMAN).//P3918 20
8
F-NT2RP3000439//HYPOTHETICAL 46.4 KD PROTEIN IN FF
H-GRPE INTERGENIC REGION.//9.8e-10:201:26//ESCHERI
CHIA COLI.//P37908
F-NT2RP3000441//PROTEIN-EXPORT MEMBRANE PROTEIN SE
CG HOMOLOG.//0.91:48:35//MYCOBACTERIUM LEPRAE.//P3
8388
F-NT2RP3000449//HOMEBOX PROTEIN HOX-B8 (CHOX-2.4)
(FRAGMENT).//1.0:42:33//GALLUS GALLUS (CHICKEN)./
/P23681
F-NT2RP3000451
F-NT2RP3000456//COLLAGEN ALPHA 1(I) CHAIN (FRAGMEN
TS).//0.00018:178:36//RATTUS NORVEGICUS (RAT).//PO
2454
F-NT2RP3000484//METALLOTHIONEIN-III (MT-III) (GROW
TH INHIBITORY FACTOR) (GIF).//0.098:40:27//BOS TAUR
US (BOVINE).//P37359
F-NT2RP3000487//SULFATED SURFACE GLYCOPROTEIN 185
(SSG 185).//0.00037:16:81//VOLVOX CARTERI.//P21997
F-NT2RP3000512 40
F-NT2RP3000526//HYPOTHETICAL NIN REGION PROTEIN OR
F56.//0.51:37:43//BACTERIOPHAGE LAMBDA.//P03769
F-NT2RP3000527//ZINC FINGER PROTEIN 83 (ZINC FINGE
R PROTEIN HPF1).//1.0e-16:234:30//HOMO SAPIENS (HU
MAN).//P51522
F-NT2RP3000531//POLIOVIRUS RECEPTOR PRECURSOR (CD1
55 ANTIGEN).//3.4e-15:192:30//HOMO SAPIENS (HUMA
N).//P15151
F-NT2RP3000542//CYTOCHROME C OXIDASE POLYPEPTIDE I
I (EC 1.9.3.1) (FRAGMENT).//0.60:51:39//ASTERINA P 50

3030

ECTINIFERA (STARFISH).//P11958
F-NT2RP3000561//HYPOTHETICAL ATP-BINDING PROTEIN M
J0423.//0.79:53:32//METHANOCOCCUS JANNASCHII.//Q57
866
F-NT2RP3000562//ACCESSORY GLAND PEPTIDE PRECURSOR
(PARAGONIAL PEPTIDE B).//0.99:26:34//DROSOPHILA MA
URITIANA (FRUIT FLY), AND DROSOPHILA SIMULANS (FRU
IT FLY).//018666
F-NT2RP3000578//HYPOTHETICAL 49.8 KD PROTEIN IN RP
L14B-GPA1 INTERGENIC REGION.//1.5e-26:127:37//SACC
HAROMYCES CEREVISIAE (BAKER'S YEAST).//P38755
F-NT2RP3000582
F-NT2RP3000584//METALLOTHIONEIN-II (MT-II).//0.28:
27:29//MUS MUSCULUS (MOUSE).//P02798
F-NT2RP3000590//UVS-2 PROTEIN.//4.8e-10:113:33//NE
UROSPORA CRASSA.//P33288
F-NT2RP3000592//TRANSCRIPTION INITIATION FACTOR TF
IID 135 KD SUBUNIT (TAFII-135) (TAFII135) (TAFII-1
30) (TAFII130).//0.00087:178:31//HOMO SAPIENS (HUM
AN).//000268
F-NT2RP3000596//YEMANUCLEIN-ALPHA.//1.8e-05:98:34/
/DROSOPHILA MELANOGASTER (FRUIT FLY).//P25992
F-NT2RP3000599//SPLICEOSOME ASSOCIATED PROTEIN 62
(SAP 62) (SF3A66).//0.00095:90:37//HOMO SAPIENS (H
UMAN).//Q15428
F-NT2RP3000603//5E5 ANTIGEN.//1.0e-09:181:34//RATT
US NORVEGICUS (RAT).//Q63003
F-NT2RP3000605//STEROL REGULATORY ELEMENT BINDING
PROTEIN-1 (SREBP-1) (STEROL REGULATORY ELEMENT-BIN
DING TRANSCRIPTION FACTOR 1).//0.00098:76:34//HOMO
SAPIENS (HUMAN).//P36956
F-NT2RP3000622//HYPOTHETICAL PROTEIN MG096 HOMOLOG
5 (P02_ORF427).//0.15:52:36//MYCOPLASMA PNEUMONIA
E.//P75277
F-NT2RP3000624//HYPOTHETICAL PROTEIN KIAA0256.//5.
4e-16:222:31//HOMO SAPIENS (HUMAN).//Q93073
F-NT2RP3000628
F-NT2RP3000632//ZINC FINGER PROTEIN 90 (ZFP-90) (Z
INC FINGER PROTEIN NK10).//2.0e-16:52:63//MUS MUSC
ULUS (MOUSE).//Q61967
F-NT2RP3000644//!!!! ALU SUBFAMILY SQ WARNING ENTR
Y !!!!//6.7e-40:102:79//HOMO SAPIENS (HUMAN).//P39
194
F-NT2RP3000661//HYPOTHETICAL 139.1 KD PROTEIN C08B
11.3 IN CHROMOSOME II.//6.0e-08:83:36//CAENORHABDI
TIS ELEGANS.//Q09441
F-NT2RP3000665//HOMEBOX PROTEIN PROPHET OF PIT-1
(PROP-1) (PITUITARY SPECIFIC HOMEODOMAIN FACTOR)./
/0.13:48:35//HOMO SAPIENS (HUMAN).//075360
F-NT2RP3000685//HYPOTHETICAL 33.5 KD PROTEIN IN CA

3031

T1 5' REGION (ORFY) //0.26:202:23//CLOSTRIDIUM KLUY VERI. //P38943
 F-NT2RP3000690//INORGANIC PYROPHOSPHATASE (EC 3.6.1.1) (PYROPHOSPHATE PHOSPHO- HYDROLASE) (PPASE) //0.99:131:26//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P00817
 F-NT2RP3000736//HYPOTHETICAL 28.7 KD PROTEIN IN RN R3-ARC15 INTERGENIC REGION. //3.5e-27:211:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST). //P40516
 F-NT2RP3000739//HYPOTHETICAL 33.5 KD PROTEIN C1D4. 10 02C IN CHROMOSOME I. //6.0e-23:114:42//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10149
 F-NT2RP3000742//1-PHOSPHATIDYLINOSITOL-4,5-BISPHOSPHATE PHOSPHODIESTERASE DELTA 1 (EC 3.1.4.11) (PLC-DELTA-1) (PHOSPHOLIPASE C-DELTA-1) (PLC-III). //6.7e-12:85:36//RATTUS NORVEGICUS (RAT). //P10688
 F-NT2RP3000753//CELL SURFACE GLYCOPROTEIN 1 PRECURSOR (OUTER LAYER PROTEIN B) (S-LAYER PROTEIN 1). //0.00011:208:28//CLOSTRIDIUM THERMOCELLUM. //Q06852
 F-NT2RP3000759//ADP-RIBOSYLATION FACTOR 6. //8.1e-2 20 8:141:38//GALLUS GALLUS (CHICKEN). //P26990
 F-NT2RP3000815//CYTOCHROME C-551 (C551) (CYTOCHROME C8). //0.24:45:37//PSEUDOMONAS DENITRIFICANS. //P0103
 F-NT2RP3000825//ALPHA-LACTALBUMIN (LACTOSE SYNTHASE B PROTEIN (EC 2.4.1.22)). //0.82:51:39//MACROPUS RUFUGRISEUS (RED-NECKED WALLABY). //P07458
 F-NT2RP3000826//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS). //0.025:79:37//BOS TAURUS (BOVINE). //P25508
 F-NT2RP3000836//HYPOTHETICAL PROTEIN IN KSGA 3' REGION (ORF L5) (FRAGMENT). //0.85:36:47//MYCOPLASMA CAPRICOLUM. //P43040
 F-NT2RP3000841//UDP-GLUCURONOSYLTRANSFERASE 1-7 PRECURSOR, MITOCHONDRIAL (EC 2.4.1.17) (UDPGT) (UGT1*7) (UGT1-07) (UGT1.7) (UGT1A7) (UGTP4) (FRAGMENT). //1.0:70:34//MUS MUSCULUS (MOUSE). //Q62452
 F-NT2RP3000845//PUTATIVE SERINE/THREONINE-PROTEIN KINASE P78 (EC 2.7.1.-). //5.2e-72:247:61//HOMO SAPIENS (HUMAN). //P27448
 F-NT2RP3000847//HYPOTHETICAL PROTEIN KIAA0161. //0. 40 037:55:30//HOMO SAPIENS (HUMAN). //P50876
 F-NT2RP3000850//!!!! ALU SUBFAMILY SQ WARNING ENTRY !!!!! //7.4e-31:90:75//HOMO SAPIENS (HUMAN). //P39194
 F-NT2RP3000852//HYDROPHOBIC SEED PROTEIN (HPS). //0.33:23:69//GLYCINE MAX(SOYBEAN). //P24337
 F-NT2RP3000859//IMMEDIATE-EARLY PROTEIN. //3.6e-07:189:25//HERPESVIRUS SAIMIRI (STRAIN 11). //Q01042
 F-NT2RP3000865
 F-NT2RP3000868//MYOSIN HEAVY CHAIN, CARDIAC MUSCLE 50

3032

ISOFORM (FRAGMENT). //1.4e-09:232:28//GALLUS GALLUS (CHICKEN). //P29616
 F-NT2RP3000869//CUTICLE COLLAGEN 2. //4.5e-08:58:46 //CAENORHABDITIS ELEGANS. //P17656
 F-NT2RP3000875//HOMEODOMAIN PROTEIN CDX-2 (CAUDAL-TYPE HOMEODOMAIN PROTEIN 2). //0.90:62:37//MUS MUSCULUS (MOUSE). //P43241
 F-NT2RP3000901//COLLAGEN ALPHA 1(I) CHAIN (FRAGMENTS). //0.99:124:33//BOSTAURUS (BOVINE). //P02453
 F-NT2RP3000904
 F-NT2RP3000917//DHP1 PROTEIN. //6.5e-60:229:55//SCHIZOSACCHAROMYCES POMBE(FISSION YEAST). //P40848
 F-NT2RP3000919//HYPOTHETICAL 33.5 KD PROTEIN C1D4. 02C IN CHROMOSOME I. //2.4e-19:159:34//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST). //Q10149
 F-NT2RP3000968//40S RIBOSOMAL PROTEIN S15A. //3.7e-48:73:98//HOMO SAPIENS(HUMAN), AND RATTUS NORVEGICUS (RAT). //P39027
 F-NT2RP3000980//COPA/INCA PROTEIN (REPA3 PROTEIN). //0.24:19:47//ESCHERICHIA COLI. //P13946
 F-NT2RP3000994//MATERNAL EFFECT PROTEIN STAUFEN. //1.4e-10:78:48//DROSOPHILA MELANOGASTER (FRUIT FLY). //P25159
 F-NT2RP3001004//HYPOTHETICAL 7.6 KD PROTEIN B0563. 8 IN CHROMOSOME X. //0.70:50:32//CAENORHABDITIS ELEGANS. //Q11084
 F-NT2RP3001007
 F-NT2RP3001055//N-TERMINAL ACETYLTRANSFERASE COMPLEX ARD1 SUBUNIT HOMOLOG. //1.3e-05:138:28//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD). //P36416
 F-NT2RP3001057//ZINC FINGER PROTEIN 45 (BRC1744). //4.0e-28:141:51//HOMOSAPIENS (HUMAN). //Q02386
 F-NT2RP3001081//HYPOTHETICAL 46.4 KD PROTEIN T16H1 2.5 IN CHROMOSOME III. //3.8e-08:144:29//CAENORHABDITIS ELEGANS. //P34568
 F-NT2RP3001084//SPIROIN 2 (DRAGLINE SILK FIBROIN 2) (FRAGMENT). //3.4e-06:217:32//NEPHILA CLAVIPES (ORB SPIDER). //P46804
 F-NT2RP3001096//SYNAPTONEMAL COMPLEX PROTEIN SC65. //1.1e-30:244:33//RATTUS NORVEGICUS (RAT). //Q64375
 F-NT2RP3001107//ARYLSULFATASE F (EC 3.1.6.-) (ASF) (FRAGMENT). //0.041:47:44//HOMO SAPIENS (HUMAN). //P54793
 F-NT2RP3001109
 F-NT2RP3001111//MALE SPECIFIC SPERM PROTEIN MST84D C. //0.17:28:39//DROSOPHILA MELANOGASTER (FRUIT FLY). //Q01644
 F-NT2RP3001113//INVOLUCRIN. //0.00036:192:23//MUS MUSCULUS (MOUSE). //P48997
 F-NT2RP3001115

3033

F-NT2RP3001116//AMINOPEPTIDASE G (EC 3.4.11.-) (FRAGMENT).//0.99:29:51//STREPTOMYCES LIVIDANS.//Q54340
 F-NT2RP3001119//COLLAGEN ALPHA 4(IV) CHAIN (FRAGMENT).//0.0015:73:39//BOS TAURUS (BOVINE).//Q29442
 F-NT2RP3001120//ZINC FINGER PROTEIN ZFP-36 (FRAGMENT).//1.3e-57:229:52//HOMO SAPIENS (HUMAN).//P16415
 F-NT2RP3001126//HYPOTHETICAL 91.2 KD PROTEIN IN RP S4B-SCH9 INTERGENIC REGION.//2.8e-07:83:34//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P38888
 F-NT2RP3001133//CALCIUM BINDING PROTEIN.//2.0e-08:171:32//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P35085
 F-NT2RP3001140//F-SPONDIN PRECURSOR.//2.0e-147:244:97//RATTUS NORVEGICUS (RAT).//P35446
 F-NT2RP3001147//TROPOMYOSIN 2 (TMII).//0.11:159:23//SCHISTOSOMA MANSONI (BLOOD FLUKE).//P42638
 F-NT2RP3001150//OCTAPEPTIDE-REPEAT PROTEIN T2.//6.2e-09:163:25//MUS MUSCULUS (MOUSE).//Q06666
 F-NT2RP3001155//DNA POLYMERASE ALPHA-BINDING PROTEIN (POB1/CTF4 PROTEIN) (CHROMOSOME REPLICATION PROTEIN CHL15).//4.1e-05:244:23//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q01454
 F-NT2RP3001176//LEUKOSIALIN PRECURSOR (LEUCOCYTE SIALOGLYCOPROTEIN) (SIALOPHORIN) (CD43) (LY 48) (B CELL DIFFERENTIATION ANTIGEN LP-3).//0.21:136:26//MUS MUSCULUS (MOUSE).//P15702
 F-NT2RP3001214//SAP1 PROTEIN.//0.058:133:30//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P39955
 F-NT2RP3001216//CYLICIN I (MULTIPLE-BAND POLYPEPTIDE 1) (FRAGMENT).//2.1e-08:137:33//HOMO SAPIENS (HUMAN).//P35663
 F-NT2RP3001221//GAMMA-BUTYROBETAINE, 2-OXOGLUTARATE DIOXYGENASE (EC 1.14.11.1) (GAMMA-BUTYROBETAINE HYDROXYLASE).//4.2e-05:131:26//PSEUDOMONAS SP. (STRAIN AK-1).//P80193
 F-NT2RP3001232//HYPOTHETICAL PROTEIN PRECURSOR IN CS5 3' REGION (FRAGMENT).//0.75:57:31//ESCHERICHIA COLI.//P33792
 F-NT2RP3001236//TRANSFORMING PROTEIN MAF.//0.017:136:30//AVIAN MUSCULOAPONEUROTIC FIBROSARCOMA VIRUS AS42.//P23091
 F-NT2RP3001239//ELECTROMOTOR NEURON-ASSOCIATED PROTEIN 1 (FRAGMENT).//4.2e-55:221:49//TORPEDO CALIFORNICA (PACIFIC ELECTRIC RAY).//P14400
 F-NT2RP3001245
 F-NT2RP3001253//TROPOMYOSIN 2, MUSCLE THORACIC ISOFORM (TROPOMYOSIN I).//0.0042:142:24//DROSOPHILA MELANOGASTER (FRUIT FLY).//P09491

3034

F-NT2RP3001260//COLLAGEN ALPHA 4(IV) CHAIN PRECURSOR.//0.0011:89:43//HOMO SAPIENS (HUMAN).//P53420
 F-NT2RP3001268//ZINC FINGER PROTEIN 45 (BRC1744).//9.0e-29:194:44//HOMOSAPIENS (HUMAN).//Q02386
 F-NT2RP3001272//HYPOTHETICAL 75.2 KD PROTEIN C13F 4.08C IN CHROMOSOME I.//8.2e-17:183:26//SCHIZOSACCHAROMYCES POMBE (FISSION YEAST).//Q10199
 F-NT2RP3001274//SERINE/THREONINE PROTEIN PHOSPHATASE 5 (EC 3.1.3.16) (PP5) (PROTEIN PHOSPHATASE T) (PPT) (FRAGMENT).//1.7e-09:78:39//MUS MUSCULUS (MOUSE).//Q60676
 F-NT2RP3001281//!!!! ALU SUBFAMILY J WARNING ENTRY !!!!!//7.7e-08:38:71//HOMO SAPIENS (HUMAN).//P39188
 F-NT2RP3001297//HYPOTHETICAL PROTEIN KIAA0281 (HA6725).//2.2e-57:159:70//HOMO SAPIENS (HUMAN).//Q92556
 F-NT2RP3001307//SPERM PROTAMINE P1.//0.21:46:39//ORNITHORHYNCHUS ANATINUS (DUCKBILL PLATYPUS).//P35307
 F-NT2RP3001318
 F-NT2RP3001325//ENHANCER OF RUDIMENTARY HOMOLOG.//1.0:73:24//BRACHYDANIORERIO (ZEBRAFISH) (ZEBRA DANIO).//Q98874
 F-NT2RP3001338//ZINC FINGER PROTEIN 29 (ZINC FINGER PROTEIN KOX26) (FRAGMENT).//0.0021:56:35//HOMO SAPIENS (HUMAN).//P17037
 F-NT2RP3001339//CITRON PROTEIN.//3.6e-06:90:33//MUS MUSCULUS (MOUSE).//P49025
 F-NT2RP3001340//HYPOTHETICAL PROTEIN UL61.//7.2e-11:202:34//HUMAN CYTOMEGALOVIRUS (STRAIN AD169).//P16818
 F-NT2RP3001355//TRICARBOXYLATE TRANSPORT PROTEIN PRECURSOR (CITRATE TRANSPORT PROTEIN) (CTP) (TRICARBOXYLATE CARRIER PROTEIN).//7.7e-16:129:33//HOMO SAPIENS (HUMAN).//P53007
 F-NT2RP3001356//RAS-RELATED PROTEIN RABA (FRAGMENT).//0.00041:66:28//DICTYOSTELIUM DISCOIDEUM (SLIME MOLD).//P34141
 F-NT2RP3001374
 F-NT2RP3001383//PTB-ASSOCIATED SPLICING FACTOR (PSF).//2.5e-06:190:32//HOMO SAPIENS (HUMAN).//P23246
 F-NT2RP3001384//CHORION PROTEIN S15.//0.00079:94:37//DROSOPHILA VIRILIS (FRUIT FLY).//P13424
 F-NT2RP3001392//VPU PROTEIN (ORF-X PROTEIN) (UPX PROTEIN).//1.0:22:45//CAPRINE ARTHRITIS ENCEPHALITIS VIRUS (CAEV).//P31834
 F-NT2RP3001396//HYPOTHETICAL 8.1 KD PROTEIN (ORF 4).//1.0:37:32//STRAWBERRY MILD YELLOW EDGE-ASSOCIATED VIRUS (SMYEA).//Q00848

3035

F-NT2RP3001398//KRUEPPEL-RELATED ZINC FINGER PROTEIN 2 (HXR2 PROTEIN) (FRAGMENT).//1.9e-08:45:37//HOMO SAPIENS (HUMAN).//P10073
 F-NT2RP3001399//SSU72 PROTEIN.//7.3e-18:84:52//SACCHAROMYCES CEREVISIAE(BAKER'S YEAST).//P53538
 F-NT2RP3001407//SCY1 PROTEIN.//1.5e-08:143:25//SACCHAROMYCES CEREVISIAE(BAKER'S YEAST).//P53009
 F-NT2RP3001420//HYPOTHETICAL 7.9 KD PROTEIN.//0.25:41:26//VACCINIA VIRUS(STRAIN COPENHAGEN).//P20542
 F-NT2RP3001426//DNAJ PROTEIN.//7.5e-15:78:43//HAEMOPHILUS INFLUENZAE.//P43735
 F-NT2RP3001427//WERNER SYNDROME HELICASE.//3.6e-13:159:33//HOMO SAPIENS(HUMAN).//Q14191
 F-NT2RP3001428//NUCLEOPROTEIN TPR.//1.8e-53:117:99//HOMO SAPIENS (HUMAN).//P12270
 F-NT2RP3001432//NADH-UBIQUINONE OXIDOREDUCTASE CHAIN 5 (EC 1.6.5.3) (FRAGMENT).//0.96:52:21//TARSUS SYRICHIA (TARSIER).//Q36151
 F-NT2RP3001447//HYPOTHETICAL 5.5 KD PROTEIN IN REPLICATION ORIGIN REGION(ORF1).//0.96:45:35//ESCHERICHIA COLI.//P14505
 F-NT2RP3001449//HOMEBOX PROTEIN SAX-1 (CHOX-3) (FRAGMENT).//0.0043:53:43//GALLUS GALLUS (CHICKEN).//P19601
 F-NT2RP3001453//MALE SPECIFIC SPERM PROTEIN MST84DB.//0.0048:65:40//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01643
 F-NT2RP3001457//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS28.//0.55:121:20//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//Q02767
 F-NT2RP3001459//MYOSIN IC HEAVY CHAIN.//0.10:126:34//ACANTHAMOEBA CASTELLANII (AMOEBA).//P10569
 F-NT2RP3001472//NONHISTONE CHROMOSOMAL PROTEIN 6A.//3.0e-14:87:43//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P11632
 F-NT2RP3001490//METALLOTHIONEIN-LIKE PROTEIN LSC54.//1.0:39:35//BRASSICANAPUS (RAPE).//P43402
 F-NT2RP3001495//UBIQUITIN--PROTEIN LIGASE RSP5 (EC 6.3.2.-).//3.3e-14:148:35//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P39940
 F-NT2RP3001497//COLLAGEN ALPHA 1(XII) CHAIN (FRAGMENTS).//0.13:44:38//BOS TAURUS (BOVINE).//P25508
 F-NT2RP3001527//SPERM PROTAMINE P1.//0.35:29:37//DIDELPHIS MARSUPIALIS VIRGINIANA (NORTH AMERICAN OP OSSUM), AND MONODELPHIS DOMESTICA (SHORT-TAILED GREY OPOSSUM).//P35305
 F-NT2RP3001529//HYPOTHETICAL 43.3 KD GTP-BINDING PROTEIN IN DACB-RPMA INTERGENIC REGION.//3.3e-21:125:37//ESCHERICHIA COLI.//P42641

3036

F-NT2RP3001538//HNF3/FH TRANSCRIPTION FACTOR GENESIS (WINGED HELIX PROTEIN CWH-3).//0.13:53:39//GALLUS GALLUS (CHICKEN).//P79772
 F-NT2RP3001554//ELECTROMOTOR NEURON-ASSOCIATED PROTEIN 2 (FRAGMENT).//2.3e-48:137:52//TORPEDO CALIFORNICA (PACIFIC ELECTRIC RAY).//P14401
 F-NT2RP3001580//GERM CELL-LESS PROTEIN.//8.2e-18:100:42//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01820
 10 F-NT2RP3001587//UBIQUITIN-ACTIVATING ENZYME E1-LIKE (POLYMERASE-INTERACTING PROTEIN 2).//2.0e-47:188:51//SACCHAROMYCES CEREVISIAE (BAKER'S YEAST).//P52488
 F-NT2RP3001589//!!!! ALU SUBFAMILY SP WARNING ENTRY !!!!!//7.4e-41:87:80//HOMO SAPIENS (HUMAN).//P39193
 F-NT2RP3001607//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.34) (AGL).//1.0:49:32//DICENTRARCHUS LABRAX (EUROPEAN SEA BASS).//Q36362
 20 F-NT2RP3001608//EXTENSIN PRECURSOR (PROLINE-RICH GLYCOPROTEIN).//0.0013:177:25//ZEA MAYS (MAIZE).//P14918
 F-NT2RP3001621//MALE SPECIFIC SPERM PROTEIN MST84D.//0.84:29:37//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q01645
 F-NT2RP3001629//RAS-RELATED C3 BOTULINUM TOXIN SUBSTRATE 1 (P21-RAC1) (FRAGMENTS).//0.91:57:24//CAVIA PORCELLUS (GUINEA PIG).//P80236
 F-NT2RP3001634//!!!! ALU SUBFAMILY SB WARNING ENTRY !!!!!//8.9e-11:73:54//HOMO SAPIENS (HUMAN).//P39189
 F-NT2RP3001642//HYPOTHETICAL PROTEIN KIAA0210.//1.1e-12:117:29//HOMO SAPIENS (HUMAN).//Q92609
 F-NT2RP3001646//HYPOTHETICAL 29.3 KD PROTEIN (ORF92).//0.0092:69:34//ORGYIA PSEUDOTSUGATA MULTICAPSID POLYHEDROSIS VIRUS (OPHNVPV).//010341
 F-NT2RP3001671//RING CANAL PROTEIN (KELCH PROTEIN).//0.0042:55:41//DROSOPHILA MELANOGASTER (FRUIT FLY).//Q04652
 40 F-NT2RP3001672
 F-NT2RP3001676//GTP-BINDING PROTEIN LEPA (FRAGMENT).//1.2e-15:56:62//PSEUDOMONAS FLUORESCENS.//P26843
 F-NT2RP3001678//SPIROIN 2 (DRAGLINE SILK FIBROIN 2) (FRAGMENT).//0.054:187:31//NEPHILA CLAVIPES (OR B SPIDER).//P46804
 F-NT2RP3001679//HYPOTHETICAL 68.7 KD PROTEIN ZK757.1 IN CHROMOSOME III.//1.5e-07:63:44//CAENORHABDITIS ELEGANS.//P34679
 50 F-NT2RP3001688//GLUCOAMYLASE S1 PRECURSOR (EC 3.2.

3037

1.3) (GLUCAN 1,4-ALPHA-GLUCOSIDASE) (1,4-ALPHA-D-G
LUCAN GLUCOHYDROLASE) (GAI).//1.0:83:28//SACCHAROM
YCES DIASTATICUS (YEAST).//P04065
F-NT2RP3001690//MYOSIN HEAVY CHAIN, CARDIAC MUSCLE
BETA ISOFORM.//0.021:247:24//HOMO SAPIENS (HUMA
N).//P12883
F-NT2RP3001698
F-NT2RP3001708//TWISTED GASTRULATION PROTEIN PRECU
RSOR.//7.7e-12:73:43//DROSOPHILA MELANOGASTER (FRU
IT FLY).//P54356
F-NT2RP3001712//CEC-1 PROTEIN.//1.9e-07:121:29//CA
ENORHABDITIS ELEGANS.//P34618
F-NT2RP3001716//SALIVARY GLUE PROTEIN SGS-3 PRECUR
SOR.//0.89:54:40//DROSOPHILA SIMULANS (FRUIT FLY).
//P13729
F-NT2RP3001724//CHROMODOMAIN-HELICASE-DNA-BINDING
PROTEIN 1 (CHD-1).//7.5e-41:164:48//HOMO SAPIENS
(HUMAN).//014646
F-NT2RP3001727//HYPOTHETICAL 37.7 KD PROTEIN ZK68
6.3 IN CHROMOSOME III.//1.5e-51:240:41//CAENORHABD
ITIS ELEGANS.//P34669
F-NT2RP3001730//SEPTIN 2 HOMOLOG (FRAGMENT).//2.4e
-122:267:86//HOMO SAPIENS (HUMAN).//Q14141
F-NT2RP3001739//INTESTINAL SODIUM/DICARBOXYLATE CO
TRANSPORTER (NA(+)/DICARBOXYLATE COTRANSPORTER).//
0.99:63:34//RATTUS NORVEGICUS (RAT).//P70545
F-NT2RP3001752//!!!! ALU SUBFAMILY SP WARNING ENTR
Y !!!!!/4.0e-21:60:85//HOMO SAPIENS (HUMAN).//P391
93
F-NT2RP3001753//HYPOTHETICAL PROTEIN KIAA0127.//7. 30
9e-12:83:44//HOMO SAPIENS (HUMAN).//Q14140
F-NT2RP3001764//DUAL SPECIFICITY PROTEIN PHOSPHATA
SE 6 (EC 3.1.3.48) (EC3.1.3.16) (DUAL SPECIFICITY
PROTEIN PHOSPHATASE PYST1).//7.7e-25:146:36//HOMO
SAPIENS (HUMAN).//Q16828
F-NT2RP3001777//SERINE/THREONINE-PROTEIN KINASE ST
E20 HOMOLOG (EC 2.7.1.-).//0.0096:204:25//CANDIDA
ALBICANS (YEAST).//Q92212
F-NT2RP3001782//ATP SYNTHASE PROTEIN 8 (EC 3.6.1.3
4) (A6L).//0.91:34:44//PONGO PYGMAEUS ABELII (SUMA 40
TRAN ORANGUTAN).//P92694
F-NT2RP3001792//HETEROGENEOUS NUCLEAR RIBONUCLEOPR
OTEIN M (HNRNP M).//1.8e-33:159:53//HOMO SAPIENS
(HUMAN).//P52272
F-NT2RP3001799//LIGHT-HARVESTING PROTEIN B800/830/
1020, ALPHA-2 CHAIN (EHS-ALPHA-2) (ANTENNA PIGMENT
PROTEIN, ALPHA-2 CHAIN).//0.14:46:28//ECTOTHORHO
DOSPIRA HALOCHLORIS.//P80103
F-NT2RP3001819//PROCOLLAGEN ALPHA 2(I) CHAIN PRECU
RSOR.//0.00030:77:36//HOMO SAPIENS (HUMAN).//P0812 50

3038

3
F-NT2RP3001844//OCTAMER-BINDING TRANSCRIPTION FACT
OR 1 (OTF-1) (NF-A1) (FRAGMENT).//0.99:43:34//MACR
OPUS EUGENII (TANMAR WALLABY).//Q28466
F-NT2RP3001854//FIBRINOGEN- AND IG-BINDING PROTEIN
PRECURSOR (MRP PROTEIN).//9.3e-10:213:24//STREPTO
COCCUS PYOGENES.//P30141
F-NT2RP3001855//HOMEODOMAIN PROTEIN PKNOX1 (HOMEODOM
AIN PROTEIN PREP-1).//2.6e-61:220:60//HOMO SAPIENS (HU
MAN).//P55347
F-NT2RP3001857//VEGETATIBLE INCOMPATIBILITY PROTEI
N HET-E-1.//1.0e-13:213:24//PODOSPORA ANSERINA.//Q
00808
F-NT2RP3001896//SPLICEOSOME ASSOCIATED PROTEIN 62
(SAP 62) (SF3A66).//0.074:124:34//HOMO SAPIENS (HU
MAN).//Q15428
F-NT2RP3001898//REGULATORY PROTEIN E2.//0.36:131:2
9//CANINE ORAL PAPILLOMAVIRUS (COPV).//Q89420
F-NT2RP3001915//CHITIN BIOSYNTHESIS PROTEIN CHS5
(CAL3 PROTEIN).//0.0021:237:23//SACCHAROMYCES CERE
VISIAE (BAKER'S YEAST).//Q12114
F-NT2RP3001926//HYPOTHETICAL 14.0 KD PROTEIN IN RP
L15B-GCR3 INTERGENIC REGION.//1.0:63:34//SACCHAROM
YCES CEREVISIAE (BAKER'S YEAST).//Q03880
F-NT2RP3001929//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/1.4e-14:35:60//HOMO SAPIENS (HUMAN).//P391
95
F-NT2RP3001931//HYPOTHETICAL 59.3 KD PROTEIN IN TA
P42-ARP9 INTERGENIC REGION.//0.86:162:24//SACCHARO
MYCES CEREVISIAE (BAKER'S YEAST).//Q05040
F-NT2RP3001938//GLYCOPROTEIN GP50.//0.0036:54:40//
PSEUDORABIES VIRUS (STRAIN RICE) (PRV).//P07645
F-NT2RP3001943//33.2 KD PROTEIN IN DIND-RPH INTERG
ENIC REGION (ORF X).//1.0:113:27//ESCHERICHIA COL
I.//P23839
F-NT2RP3001944//HYPOTHETICAL 47.6 KD PROTEIN C16C1
0.5 IN CHROMOSOME III.//4.1e-56:208:47//CAENORHABD
ITIS ELEGANS.//Q09251
F-NT2RP3001969//PUFF II/9-2 PROTEIN PRECURSOR.//0.
0078:149:26//SCIARA COPROPHILA (FUNGUS GNAT).//P22
312
F-NT2RP3001989//SPERM PROTAMINE P1 (CYSTEINE-RICH
PROTAMINE).//1.0:41:31//MUS MUSCULUS (MOUSE).//P02
319
F-NT2RP3002002//!!!! ALU SUBFAMILY SX WARNING ENTR
Y !!!!!/1.2e-44:69:79//HOMO SAPIENS (HUMAN).//P391
95
F-NT2RP3002004//TRANSCRIPTION FACTOR BF-2 (BRAIN F
ACTOR 2) (BF2).//0.00024:45:40//MUS MUSCULUS (MOUS
E).//Q61345